

ANSWERS TO PROBLEMS

Chapter 1 Answers

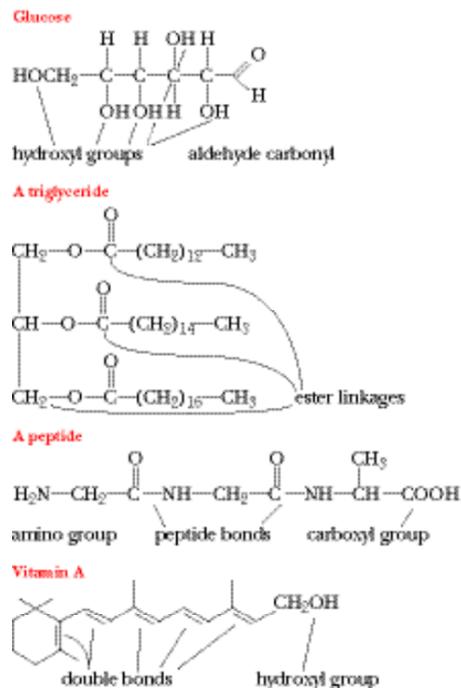
1.1 What Are the Basic Themes for This Text?

1. A polymer is a very large molecule formed by linking smaller units (monomers) together. A protein is a polymer formed by linking amino acids together. A nucleic acid is a polymer formed by linking nucleotides together. Catalysis is the process that increases the rate of chemical reactions compared to the uncatalyzed reaction. Biological catalysts are proteins in almost all cases; the only exceptions are a few types of RNA, which can catalyze some of the reactions of their own metabolism. The genetic code is the means by which the information for the structure and function of all living things is passed from one generation to the next. The sequence of purines and pyrimidines in DNA carries the genetic code (RNA is the coding material in some viruses).

1.2 What Is the Chemical Nature of Important Biomolecules?

2. The correct match of functional groups and compounds containing that functional group is given in the following list.
Amino group $\text{CH}_3\text{CH}_2\text{NH}_2$
Carbonyl group (ketone) CH_3COCH_3
Hydroxyl group CH_3OH Carboxyl group CH_3COOH Carbonyl group (aldehyde) $\text{CH}_3\text{CH}_2\text{CHO}$ Thiol group CH_3SH Ester linkage $\text{CH}_3\text{COOCH}_2\text{CH}_3$ Double bond $\text{CH}_3\text{CH}=\text{CH}_2$ Amide linkage $\text{CH}_3\text{CON}(\text{CH}_3)_2$ Ether $\text{CH}_3\text{CH}_2\text{OCH}_2\text{CH}_3$

3. The functional groups in the compounds follow:



4. Before 1828, the concept of vitalism held that organic compounds

could only be made by living systems and were beyond the realm of laboratory investigations. Wohler's synthesis showed that organic compounds, like inorganic ones, did not require a vitalistic explanation, but rather they obeyed the laws of chemistry and physics and thus were subject to laboratory investigation. Subsequently, the concept was extended to the much more complex, but still testable, discipline of biochemistry.

5. Urea, like all organic compounds, has the same molecular structure whether it is produced by a living organism or not.
- 6.



7. Five; 7 if the two cyclopropane derivatives are allowed.
8. Thirteen different alcohols, 11 aldehydes/ketones, and 10 each epoxides and ethers.

1.3 What Can Biochemistry Say About Possible Origins of Life?

9. It is generally believed that carbon is the likely basis for all life forms, earth-bound or extraterrestrial.
10. Eighteen residues would give 20^{18} or 2.6×10^{23} possibilities. Thus 19 residues would be necessary to have at least Avogadro's number (6.022×10^{23}) of possibilities.
11. The number is 4^{40} or 1.2×10^{24} , which is twice Avogadro's number.
12. RNA is capable of both coding and catalysis.
13. Catalysis allows living organisms to carry out chemical reactions much more efficiently than without catalysts.
14. Two of the most obvious advantages are speed and specificity; they also work at constant temperature or produce little heat.
15. Coding allows for reproduction of cells.
16. With respect to coding, RNA-type polynucleotides have been produced from monomers in the absence of either a preexisting RNA to be copied or an enzyme to catalyze the process. The observation that some existing RNA molecules can catalyze their own processing suggests a role for RNA in catalysis. With this dual role, RNA may have been the original informational macromolecule in the origin of life.
17. It is unlikely that cells could have arisen as bare cytoplasm without a plasma membrane. The presence of the membrane protects cellular components from the environment and prevents them from diffusing away from each other. The molecules within a cell can react more easily if they are closer to each other.

1.4 How Do Prokaryotes and Eukaryotes Differ in Levels of Organization?

18. Five differences between prokaryotes and eukaryotes are: (1) Prokaryotes do not have a well-defined nucleus, but eukaryotes have a nucleus marked off from the rest of the cell by a double membrane. (2) Prokaryotes have only a plasma (cell) membrane; eukaryotes have an extensive internal membrane system. (3)

Eukaryotic cells contain membrane-bounded organelles, while prokaryotic cells do not, (4) Eukaryotic cells are normally larger than those of prokaryotes, (5) Prokaryotes are single-celled organisms, while eukaryotes can be multicellular as well as single-celled.

19. Protein synthesis takes place on ribosomes in both prokaryotes and eukaryotes. In eukaryotes, ribosomes may be bound to the endoplasmic reticulum or found free in the cytoplasm; in prokaryotes, ribosomes are only found free in the cytoplasm.

1.5 What Are the Main Structural Features of Prokaryotic Cells?

20. It is unlikely that mitochondria would be found in bacteria. These eukaryotic organelles are enclosed by a double membrane, and bacteria do not have an internal membrane system. The mitochondria found in eukaryotic cells are about the same size as most bacteria.

1.6 What Are the Main Structural Features of Eukaryotic Cells?

21. See Section 1.6 for the functions of the parts of an animal cell, which are shown in Figure 1.10(a).
22. See Section 1.6 for the functions of the parts of a plant cell, which are shown in Figure 1.10(b).
23. In green plants photosynthesis takes place in the membrane system of chloroplasts, which are large membrane-enclosed organelles. In photosynthetic bacteria, there are extensions of the plasma membrane into the interior of the cell called chromatophores, which are the sites of photosynthesis.
24. Nuclei, mitochondria, and chloroplasts are bounded by a double membrane.
25. Nuclei, mitochondria, and chloroplasts all contain DNA. The DNA found in mitochondria and in chloroplasts differs from that in the nucleus.
26. Mitochondria carry out a high percentage of the oxidation energy-releasing reactions of the cell. They are the primary sites of ATP synthesis.
27. The Golgi apparatus is involved in binding carbohydrates to proteins and in exporting substances from the cell. Lysosomes contain hydrolytic enzymes, peroxisomes contain catalase (needed for the metabolism of peroxides), and glyoxysomes contain enzymes needed by plants for the glyoxylate cycle. All these organelles have the appearance of flattened sacs bounded by a single membrane.

1.7 How Do We Classify Organisms: Five Kingdoms or Three Domains?

28. Monera include bacteria (e. g., *E. coli*) and cyanobacteria. Protista include Euglena, Volvox, Amoeba, and Paramecium. Fungi include molds and mushrooms. Plantae include clubmosses and oak trees. Animalia include spiders, salmon, rattlesnakes, robins, and dogs.
29. The kingdom Monera consists of prokaryotes. The other four

kingdoms consist of eukaryotes.

30. The kingdom Monera is divided into the domains Eubacteria and Archaea on the basis of biochemical differences. The domain Eukarya consists of the four kingdoms of eukaryotic organisms.

1.8 Is There Common Ground for All Cells?

31. The major advantage is that of having compartments (organelles) with specialized functions (and thus division of labor). Another advantage is that cells can be much larger without surface area to volume considerations being critical because of compartmentalization.
32. See the discussion of the endosymbiotic theory in Section 1.8.
33. See Exercise 32. The division of labor in cells gives rise to greater efficiency and a larger number of individuals. This in turn allows more opportunity for evolution and speciation.

1.9 How Do Cells Use Energy?

34. Processes that release energy are favored.

1.10 What Is the Connection between Energy and Change?

35. The term "spontaneous" means energetically favored. It does not necessarily mean fast.

1.11 What Is the Criterion for Spontaneity in Biochemical Reactions?

36. The system is the nonpolar solute and water, which become more disordered when a solution is formed; ΔS_{sys} is positive but comparatively small. The ΔS_{sur} is negative and comparatively large because it is a reflection of the unfavorable enthalpy change for forming the solution (ΔH_{sys}).
37. Processes (a) and (b) are spontaneous, whereas processes (c) and (d) are not. The spontaneous processes represent an increase in disorder (increase in the entropy of the universe) and have a negative ΔG° at constant temperature and pressure, while the opposite is true of the nonspontaneous processes.
38. In all cases, there is an increase in entropy, and the final state has more possible random arrangements than the initial state.
39. Since the equation involves multiplication of ΔS by T , the value of ΔG is temperature-dependent.
40. If one considers entropy to be a measure of dispersion of energy, then at higher temperatures, it is logical that molecules would have more possible arrangements due to increased molecular motion.
41. Assuming the value of ΔS is positive, an increase in temperature will increase the - contribution of the entropy component to the overall energy change.
42. The heat exchange, getting colder, reflects only the enthalpy or ΔH component of the total energy change. The entropy change must be high enough to offset the enthalpy component and add up to an overall $-\Delta G$.
43. Entropy would increase. There are more ways that two molecules, ADP and P_i , can be randomized than a single molecule, ATP.

1.12 What Is the Connection between Thermodynamics and Life?

44. The lowering of entropy needed to give rise to organelles leads to higher entropy in the surroundings, thus increasing the entropy of the universe as a whole
45. Compartmentalization in organelles brings components of reactions into proximity with one another. The energy change of the reaction is not affected, but the availability of components allows in to proceed more readily.
46. DNA would have higher entropy with the strands separated. There are two single strands instead of one double strand, and the single strand have more conformational mobility.
47. See the answer to exercise 43. It is still unlikely that cells could have arisen as bare cytoplasm, but the question of proximity of reactants is more to the point here than the energy change of a given reaction.
48. It would be unlikely that cells of the kind we know would have evolved on a gas giant. The lack of solids and liquids on which aggregates could form would make a large difference.
49. The available materials differ from those that would have been found on earth, and conditions of temperature and pressure are very different.
50. Mars, because of conditions more like those on earth.
51. A number of energetically favorable interactions drive the process of protein folding, ultimately increasing the entropy of the universe.
52. Photosynthesis is endergonic, requiring light energy from the sun. The complete aerobic oxidation of glucose is exergonic and is a source of energy for many organisms, including humans. It would be reasonable to expect the two processes to take place differently in order to provide energy for the endergonic one.

Chapter 2 Answers

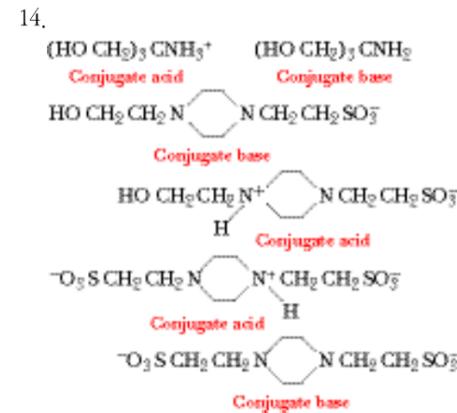
2.2 What Is a Hydrogen Bond?

1. Proteins and nucleic acids have hydrogen bonds as an important part of their structures.
2. Replication of DNA and its transcription to RNA requires hydrogen bonding of complementary bases to the DNA template strand.
3. The C—H bond is not sufficiently polar for greatly unequal distribution of electrons at its two ends. Also, there are no unshared pairs of electrons to serve as hydrogen bond acceptors.
4. Many molecules can form hydrogen bonds. Examples might be H₂O, CH₃OH, NH₃.
5. For a bond to be called a hydrogen bond, it must have a hydrogen covalently bonded to O, N, or F. This hydrogen then forms a hydrogen bond with another O, N, or F.
6. In a hydrogen-bonded dimer of acetic acid the —OH portion of the carboxyl group on molecule 1 is hydrogen-bonded to the =O portion of the carboxyl group on molecule 2, and vice versa.
7. Glucose = 17 and sorbitol = 18, ribitol + 15; each alcohol group can bond to 3 waters and the ring oxygen binds to two. The sugar alcohols bind more than the corresponding sugars.)

8. Hypoventilation decreases the pH of blood. (See BC on p. 54)
9. Aspirin is electrically neutral at the pH of the stomach and can pass the membrane more easily there than in the small intestine.
10. Positively charged ions will bind to nucleic acids as a result of electrostatic attraction to the negatively charged phosphate groups.
11. The unique fitness of water for forming hydrogen bonds determines the properties of many important biomolecules. Water can also act as an acid and a base, giving it great versatility in biochemical reactions.
12. If atoms did not differ in electronegativity, there would be no polar bonds. This would drastically affect all reactions that involve functional groups containing oxygen or nitrogen. That is most of biochemistry.

2.3 What Are Acids and Bases?

13. (CH₃)₃NH⁺ (conjugate acid)
(CH₃)₃N (conjugate base)
⁺H₃N—CH₂—COOH (conjugate acid)
⁺H₃N—CH₂—COO⁻ (conjugate base)
⁺H₃N—CH₂—COO⁻ (conjugate acid)
H₂N—CH₂—COO⁻ (conjugate base)
⁻OOC—CH₂—COOH (conjugate acid)
⁻OOC—CH₂—COO⁻ (conjugate base)
⁻OOC—CH₂—COOH (conjugate base)
HOOC—CH₂—COOH (conjugate acid)



15.
 - (a) The numerical constant equal to the concentration of the products of the dissociation divided by the concentration of the undissociated acid form: $([H^+][A^-])/[HA]$
 - (b) The qualitative or quantitative description of how much acid (HA) dissociates to hydrogen ion.
 - (c) The property of a molecule that has both a polar region and a nonpolar region.
 - (d) The amount of acid or base that can be added to a buffer before experiencing a sharp pH change.
 - (e) The point in a titration curve where the added acid or base equals the amount of buffer originally present.
 - (f) The property of a molecule that is readily soluble in water (i.e.

water loving)

- (g) The property of a molecule that is insoluble in water (i.e. water hating).
- (h) The property of a molecule that is not soluble in water. The property of a covalent bond where there is even sharing of electrons and no dipole moments (partial charges).
- (i) The property of a molecule that is soluble in water. The property of a covalent bond where the electrons are not shared evenly and dipole moments (partial charges) exist.
- (j) An experiment whereby acid or base is added stepwise to a compound and the pH is measured as a function of the added substance.

2.4 What is pH, and What Does It Have To Do with the Properties of Water?

16. The definition of pH is $-\log[H^+]$. Due to the log function, a change in concentration of 10 will lead to a change in pH of 1. The log of 10 is one. The log of 100 is 2, etc.
17. Blood plasma, pH 7.4 $[H^+] = 4.0 \times 10^{-8}$ M
Orange juice, pH 3.5 $[H^+] = 3.2 \times 10^{-4}$ M
Human urine, pH 6.2 $[H^+] = 6.3 \times 10^{-7}$ M
Household ammonia, pH 11.5 $[H^+] = 3.2 \times 10^{-12}$ M
Gastric juice, pH 1.8 $[H^+] = 1.6 \times 10^{-2}$ M
18.
 - Saliva, pH 6.5 $[H^+] = 3.2 \times 10^{-7}$ M
 - Intracellular fluid (liver), pH 6.9 $[H^+] = 1.6 \times 10^{-7}$ M
 - Tomato juice, pH 4.3 $[H^+] = 5.0 \times 10^{-5}$ M
 - Grapefruit juice, pH 3.2 $[H^+] = 6.3 \times 10^{-4}$ M
19.
 - Saliva, pH 6.5 $[OH^-] = 3.2 \times 10^{-8}$ M
 - Intracellular fluid (liver), pH 6.9 $[OH^-] = 7.9 \times 10^{-8}$ M
 - Tomato juice, pH 4.3 $[OH^-] = 2.0 \times 10^{-10}$ M
 - Grapefruit juice, pH 3.2 $[OH^-] = 1.6 \times 10^{-11}$ M

2.6 What Are Buffers, and Why Are They Important?

20. The pK of the buffer should be close to the desired buffer pH, and the substance chosen should not interfere with the reaction being studied.
21. The useful pH range of a buffer is one pH unit above and below its pKa.
22. Use the Henderson-Hasselbalch equation

$$pH = pK a^+ \log \left(\frac{[CH_3COO^-]}{[CH_3COOH]} \right)$$

$$5,00 = 4,76 + \log \left(\frac{[CH_3COO^-]}{[CH_3COOH]} \right)$$

$$0,24 = \log \left(\frac{[CH_3COO^-]}{[CH_3COOH]} \right)$$

$$[CH_3COO^-] = \text{inverse log of } 0,24 = 1,7$$

$$[CH_3COOH] \quad \quad \quad 1$$
23. Use the Henderson-Hasselbalch equation

$$pH = pK a^+ \log \left(\frac{[CH_3COO^-]}{[CH_3COOH]} \right)$$

$$4,00 = 4,76 + \log \left(\frac{[CH_3COO^-]}{[CH_3COOH]} \right)$$

$$-0,76 = \log \left(\frac{[CH_3COO^-]}{[CH_3COOH]} \right)$$

$$[CH_3COO^-] = \text{inverse log of } -0,76 = 0,17$$

$$[CH_3COOH] \quad \quad \quad 1$$

24. Use the Henderson-Hasselbalch equation

$$pH = pK a^+ \log \left(\frac{[TRIS]}{[TRIS-H^+]}\right)$$

$$8,7 = 8,3 + \log \left(\frac{[TRIS]}{[TRIS-H^+]}\right)$$

$$0,4 = \log \left(\frac{[TRIS]}{[TRIS-H^+]}\right)$$

$$[TRIS] = \text{inverse log of } 0,4 = 2,5$$

$$[TRIS-H^+] \quad \quad \quad 1$$
25. Use the Henderson-Hasselbalch equation

$$pH = pK a^+ \log \left(\frac{[HEPES]}{[HEPES-H^+]}\right)$$

$$7,9 = 7,55 + \log \left(\frac{[HEPES]}{[HEPES-H^+]}\right)$$

$$0,35 = \log \left(\frac{[HEPES]}{[HEPES-H^+]}\right)$$

$$[HEPES] = \text{inverse log of } 0,35 = 2,2$$

$$[HEPES-H^+] \quad \quad \quad 1$$

26. At pH 7.5, the ratio of $[HPO_4^{2-}]/[H_2PO_4^-]$ is 2/1 (pKa of H₂PO₄ = 7.2), as calculated using the Henderson-Hasselbalch equation. K₂HPO₄ is a source of the base form, and HCl must be added to convert one-third of it to the acid form, according to the 2/1 base/acid ratio. Weigh out 8.7 grams of K₂HPO₄ (0.05 moles, based on a formula weight of 174 grams/mole), dissolve in a small quantity of distilled water, add 16.7 mL of 1 M HCl (gives 1/3 of 0.05 moles of hydrogen ion, which converts 1/3 of the 0.05 moles of HPO₄²⁻ to H₂PO₄⁻) and dilute the resulting mixture to one liter.
27. A 2/1 ratio of the base form to acid form is still needed, because the pH of the buffer is the same in both problems. NaH₂PO₄ is a source of the acid form, and NaOH must be added to convert two thirds of it to the base form. Weigh out 6.0 grams of NaH₂PO₄ (0.05 moles, based on a formula weight of 120 grams/mole), dissolve in a small quantity of distilled water, add 33.3 mL of 1 M NaOH (gives 2/3 of 0.05 moles of hydroxide ion, which converts 2/3 of the 0.05 moles of H₂PO₄⁻ to HPO₄²⁻) and dilute the resulting mixture to one liter.
28. After mixing the buffer solution (100 mL) contains 0.75 M lactic acid and 0.25 M sodium lactate. The pKa of lactic acid is 3.86. Use the Henderson-Hasselbalch equation

$$pH = pK a^+ \log \left(\frac{[CH_3CHOHCOO^-]}{[CH_3CHOHCOOH]} \right)$$

$$pH = 3,86 + \log \left(\frac{[CH_3CHOHCOO^-]}{[CH_3CHOHCOOH]} \right)$$

$$pH = 3,86 + \log (0,25 \text{ M}/0,75 \text{ M})$$

- pH = 3,86 + (-0,48)
pH = 3,38
29. After mixing the buffer solution (100 mL) contains 0,25 M lactic acid and 0,75 M sodium lactate, The pKa of lactic acid is 3,86, Use the Henderson-Hasselbalch equation

$$\text{pH} = \text{pK} a^+ \log \left(\frac{[(\text{CH}_3\text{CHOHCOO}^-)]}{[(\text{CH}_3\text{CHOHCOOH})]} \right)$$

$$\text{pH} = 3,86 + \log \left(\frac{[(\text{CH}_3\text{CHOHCOO}^-)]}{[(\text{CH}_3\text{CHOHCOOH})]} \right)$$

$$\text{pH} = 3,86 + \log (0,75 \text{ M}/0,25 \text{ M})$$

$$\text{pH} = 3,86 + (0,48)$$

$$\text{pH} = 4,34$$
30. Use the Henderson-Hasselbalch equation

$$\text{pH} = \text{pK} a^+ \log \left(\frac{[(\text{CH}_3\text{COO}^-)]}{[(\text{CH}_3\text{COOH})]} \right)$$

$$\text{pH} = 4,76 + \log \left(\frac{[(\text{CH}_3\text{COO}^-)]}{[(\text{CH}_3\text{COOH})]} \right)$$

$$\text{pH} = 4,76 + \log \left(\frac{0,25 \text{ M}}{0,10 \text{ M}} \right)$$

$$\text{pH} = 4,76 + 0,40$$

$$\text{pH} = 5,16$$
31. Yes, it is correct, calculate the molar amounts of the two forms and insert into the Henderson-Hasselbalch equation, 2,02 g = 0,0167 mol and 5,25 g = 0,0333 mol
32. The solution is a buffer because it contains equal concentrations of TRIS in the acid and free amine forms, When the two solutions are mixed, the concentrations of the resulting solution (in the absence of reaction) are 0,05 M HCl and 0,1 M TRIS because of dilution. The HCl reacts with half the TRIS present, giving 0,05 M TRIS (protonated form) and 0,05 M TRIS (free amine form).
33. Any buffer that has equal concentrations of the acid and basic forms will have a pH equal to its pKa. Therefore the buffer from exercise 32 will have a pH of 8,3.
34. First calculate the moles of buffer that you have: 100 mL = 0,1 liters, and 0,1 liters of 0,1 M TRIS buffer is 0,01 moles. Since the buffer is at its pKa, there are equal concentrations of the acid and basic form, so the amount of TRIS is 0,005 moles, and the amount of TRIS-H⁺ is 0,005 moles, If you then add 3 mL of 1 M HCl, you will be adding 0,003 moles of H⁺. This will react as shown:

$$\text{TRIS} + \text{H}^+ \rightarrow \text{TRIS-H}^+$$
until you run out of something, which will be the H⁺ since it is the limiting reagent. The new amounts can be calculated as shown below:

$$\text{TRIS-H}^+ = 0,005 \text{ moles} + 0,003 \text{ moles} = 0,008 \text{ moles}$$

$$\text{TRIS} = 0,005 \text{ moles} - 0,003 \text{ moles} = 0,002 \text{ moles}$$
Now plug these values into the Henderson-Hasselbalch equation:

$$\text{pH} = 8,3 + \log \left(\frac{[\text{TRIS}]}{[\text{TRIS-H}^+]} \right) = 8,3 + \log(0,002/0,008)$$

$$\text{pH} = 7,70$$
35. First calculate the moles of buffer that you have: (we are going to do some rounding off) 100 mL = 0,1 liters, and 0,1 liters of 0,1 M TRIS buffer is 0,01 moles. Since the buffer is at pH 7,70, we saw in

- exercise 25 that the amount of TRIS is 0,002 moles, and the amount of TRIS-H⁺ is 0,008 moles, If you then add 3 mL of 1 M HCl, you will be adding 0,003 moles of H⁺. This will react as shown:

$$\text{TRIS} + \text{H}^+ \rightarrow \text{TRIS-H}^+$$
until you run out of something, which will be the TRIS since it is the limiting reagent. All the TRIS is converted to TRIS-H⁺:

$$\text{TRIS-H}^+ = 0,01 \text{ moles}$$

$$\text{TRIS} = \sim 0 \text{ moles}$$
We have used up the buffer capacity of the TRIS. We now have 0,003 moles of H⁺ in approximately 0,1 liters of solution. This is approximately 0,3M H⁺.

$$\text{pH} = -\log 0,3$$

$$\text{pH} = 0,52$$
36. [H⁺] = [A⁻] for pure acid, thus Ka = [H⁺]²/[HA]
[H⁺]² = Ka x [HA] -2 log [H⁺] = pKa - log [HA]

$$\text{pH} = 1/2 \{ \text{pKa} - \log [\text{HA}] \}$$
37. Use the Henderson-Hasselbalch equation, [Acetate ion]/[acetic acid] = 2,3/1
38. A substance with a pKa of 3,9 has a buffer range of 2,9 to 4,9. It will not buffer effectively at pH 7,5.
39. Use the Henderson-Hasselbalch equation, The ratio of A⁻/HA would be 3981 to 1.
40. In all cases the suitable buffer range covers a pH range of pKa +/- 1 pH units.
Lactic acid (pKa = 3,86) and its sodium salt, pH 2,86-4,86
Acetic acid (pKa = 4,76) and its sodium salt, pH 3,76-5,76
TRIS (see Table 3,4, pKa, = 8,3) in its protonated form and its free amine form, pH 7,3-9,3
HEPES (see Table 3,4, pKa, = 7,55) in its zwitterionic and its anionic form, pH 6,55-8,55
41. Several of the buffers would be suitable, namely TES, HEPES, MOPS, and PIPES, however the best buffer would be MOPS as its pKa of 7,2 is closest to the desired pH of 7,3.
42. The solution is called 0,0500 molar, even though the concentration of neither the free base nor the conjugate acid is 0,0500 M. Why is 0,0500M the correct concentration to report? Buffer concentrations are typically reported to be the sum of the two ionic forms.)
43. At the equivalence point of the titration a small amount of acetic acid remains because of the equilibrium $\text{CH}_3\text{COOH} \rightarrow \text{H}^+ + \text{CH}_3\text{COO}^-$. There is a small, but nonzero, amount of acetic acid left.
44. Buffering capacity is based upon the amounts of the acid and base forms present in the buffer solution. A solution with a high buffering capacity can react with large amount of added acid or base without drastic changes in pH. A solution with a low buffering capacity can react with only comparatively small amounts of acid or base before showing changes in pH. The more concentrated the buffer, the higher is its buffering capacity. The first buffer listed here has 10 times less buffering capacity than the second,

which in turn has 10 times less buffering capacity than the third. All three buffers have the same pH, since they all have the same relative amounts of the acid and base form.

45. It would be more effective to start with the HEPES base. You want a buffer at a pH above the pKa, which means that the base form will predominate when you have finished preparing it. It is easier to convert some of the base form to the acid form than most of the acid form to the base form.
46. In a buffer with the pH above the pKa, the base form predominates. This would be useful as a buffer for a reaction that produces H⁺ because there will be plenty of the base form to react with the hydrogen ion produced.
47. Zwitterions tend not to interfere with biochemical reactions.
48. It is useful to have a buffer that will maintain a stable pH even if assay conditions change. Dilution is one such possible change.
49. It is useful to have a buffer that will maintain a stable pH even if assay conditions change. Temperature variation is one such possible change.
50. The only zwitterion is ⁺H₃N-CH₂-COO⁻.

Chapter 3 Answers

3,1 What Are Amino Acids, and What is Their Three-Dimensional Structure?

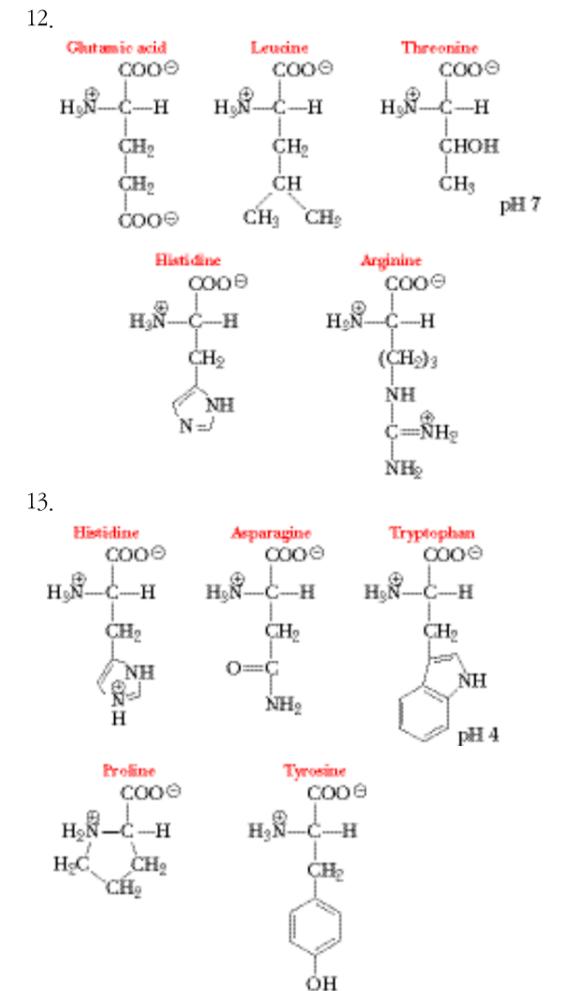
1. D- and L-amino acids have different stereochemistry around the α-carbon. Peptides that contain D-amino acids are found in bacterial cell walls and in some antibiotics.

3,2 What Are the Structures and Properties of the Individual Amino Acids?

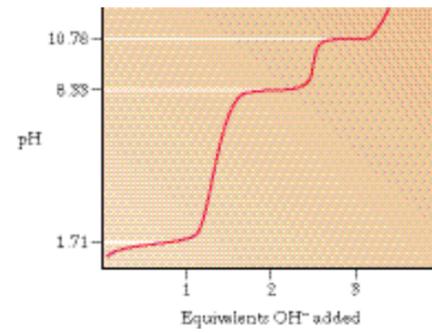
2. Proline is technically not an amino acid. Glycine contains no chiral carbon atoms.
3. An amino acid in which the R group contains the following: a hydroxyl group (serine, threonine, tyrosine) a sulfur atom (cysteine, methionine) a second chiral carbon atom (isoleucine, threonine) an amino group (lysine) an amide group (asparagine, glutamine) an acid group (aspartate, glutamate) an aromatic ring (phenylalanine, tyrosine, tryptophan) a branched side chain (leucine, valine)
4. In the peptide, Val-Met-Ser-Ile-Phe-Arg-Cys-Tyr-Leu, the polar amino acids are Ser, Arg, Cys, and Tyr; the aromatic amino acids are Phe and Tyr; and the sulfur-containing amino acids are Met and Cys.
5. In the peptide, Glu-Thr-Val-Asp-Ile-Ser-Ala, the nonpolar amino acids are Val, Ile, and Ala; the acidic amino acids are Glu and Asp.
6. Amino acids other than the usual 20 are produced by modification of one of the common amino acids. See Figure 3,5 for the structures of some modified amino acids. Hydroxyproline and hydroxylysine are found in collagen; thyroxine is found in thyroglobulin.
7. Tyrosine, tryptophan, and their derivatives.
8. A monoamine oxidase is an enzyme that degrades compounds with an amino group, including neurotransmitters; consequently, they can control a person's mental state.

9. The high concentration of tryptophan in milk protein may mildly elevate the levels of serotonin, which relaxes the brain.
10. The tryptophan in milk might make you sleepy, whereas the tyramine in cheese should pep you up.
11. The amino acids thyroxine and hydroxyproline occur in very few proteins. The genetic code does not include them, so they are produced by modification of tyrosine and proline, respectively.

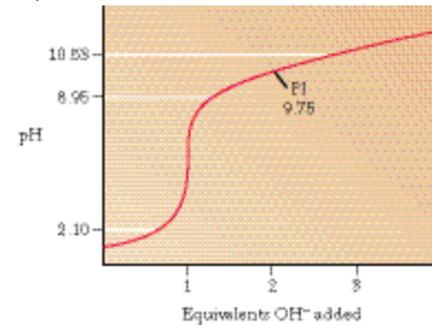
3,3 Do Amino Acids Have Specific Acid-Base Properties?



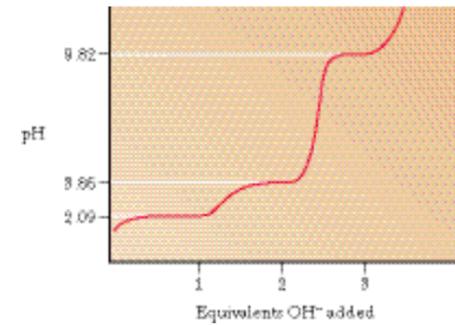
14. Histidine: imidazole is deprotonated, -amino group is predominantly deprotonated, Asparagine: -amino group is deprotonated, Tryptophan: -amino group is predominantly deprotonated, Proline: -amino group is partially deprotonated, Tyrosine: -amino group is predominantly deprotonated, phenolic hydroxyl is approximately a 50 · 50 mixture of protonated and deprotonated forms.
15. Isoelectric point for glutamic acid, 3,25; serine, 5,7; histidine, 7,58; lysine, 9,75; tyrosine, 5,65; arginine, 10,75.
16. Cysteine will have no net charge at pH 5,02 = (1,71 + 8,33)/2 (see titration curve below).



17.

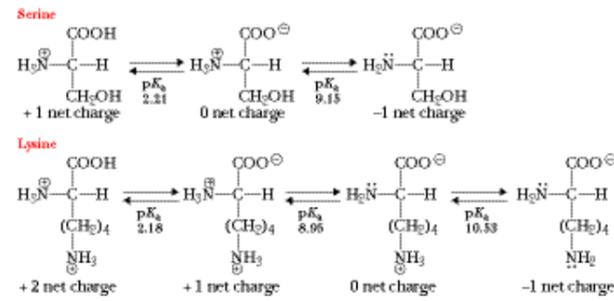
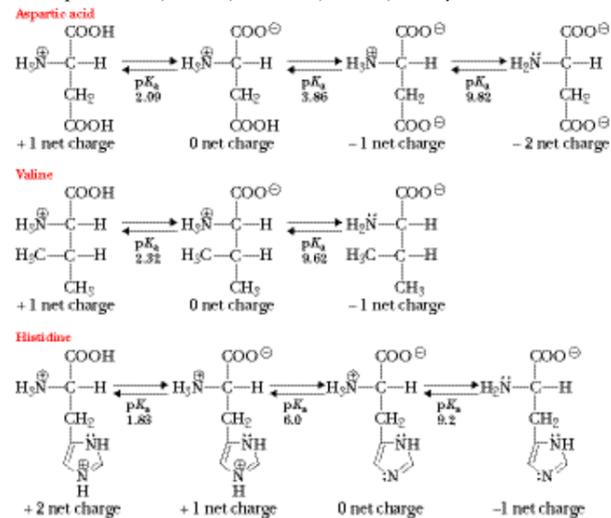


18. The conjugate acid - base pair will act as a buffer in the pH range 1.09 - 3.09.



19. They have a net charge at pH extremes, and the molecules tend to repel each other. When the molecular charge is zero, the amino acids can aggregate more easily.

20. The ionic dissociation reactions of the following amino acids: aspartic acid, valine, histidine, serine, and lysine:



21. The pKa for the ionization of the thiol group of cysteine is 8.33, so this amino acid could serve as a buffer in the -SH and -S- forms over the pH range 7.33-9.33. The - amino groups of asparagine and lysine have pKa values of 8.80 and 8.95, respectively; these are also possible buffers, but they are both near the end of their buffer ranges.

22. At pH 4, the -carboxyl group is deprotonated to a carboxylate, the side chain carboxyl is still more than 50% protonated, and both amino groups are protonated. At pH 7, both the -carboxyl group and the side chain carboxyl group are deprotonated to a carboxylate, and both amino groups are protonated. At pH 10, both the -carboxyl group and the side chain carboxyl group are deprotonated to a carboxylate, one of the amino groups is primarily deprotonated, and the other amino group is a mixture of the protonated and deprotonated forms.

23. The pI refers to the form in which both carboxyl groups are deprotonated and both amino groups protonated at pH 6.96.

23. The pI refers to the form in which both carboxyl groups are deprotonated and both amino groups protonated at pH 6.96.

24. At pH 1 the charged groups are the N-terminal NH_3^+ on valine and the protonated guanidino group on arginine, giving zero net charge. The charged groups at pH 7 are the same as pH 1 with the addition of the carboxylate group on the C-terminal leucine, giving a net charge of -1.

25. Both peptides, Phe-Glu-Ser-Met and Val-Trp-Cys-Leu, have a charge of -1 at pH 1 because of the protonated N-terminal amino group. At pH 7, the peptide on the right has no net charge because of the protonated N-terminal amino group and the ionized C-terminal carboxylate negative charge. The peptide on the left has a net charge of -1 at pH 7 because of the side-chain carboxylate group on the glutamate in addition to the charges on the N-terminal and C-terminal groups.

26. (a) Lysine because of the side chain amino group; (b) Serine because of the lack of a side chain carboxyl; (c) Histidine, because of the presence of a titratable group in the side chain; (d) Aspartate because of the carboxyl group in the side chain.

27. Glycine is frequently used as the basis of a buffer in the acid range near the pK of its carboxyl group. The useful buffer range is from pH 1.3 to 3.3.

3.4 What Is the Peptide Bond?

28. See Figure 3.9.

29. The resonance structures contribute to the planar arrangement by

giving the C-N bond partial double bond character.

30. In all cases, the yield is 0.95n. For 10 residues, that means 60% yield; for 50 residues, 8%; and for 100 residues, 0.6%. These are not satisfactory yields. Enzyme specificity gets around the problem.

31. The two peptides differ in amino acid sequence but not in composition.

32. The titration curves of the two peptides will have the same general shape. The pKa values of the -amino and -carboxyl groups will differ. Very careful work will show slight differences in side chain pKa values because of the different distances to the charged groups at the ends of the peptide. Such changes are particularly marked in proteins.

33. Asp-Leu-Phe Leu-Asp-Phe Phe-Asp-Leu Asp-Phe-Leu Leu-Phe-Asp Phe-Leu-Asp

34. DLF LDF FDL DFL LFD FLD

35. You would get $20^{100} \sim 1.27 \times 10^{130}$ molecules, which is about 10^{84} Earth volumes. The same calculation for a pentapeptide gives more comprehensible results.

36. They are relatively stable because they are zwitterions. They typically have high melting points.

37. With very little doubt, no. Compare predicting the properties of water from those of hydrogen and oxygen, in either atomic or molecular form. If you knew the properties of the protein, you might be able to do the reverse to some extent.

38. These two peptides differ chemically. The open chain has a free C-terminal and N-terminal, but the cyclic peptide has only peptide bonds.

39. Both the C-terminal and the N-terminals of the open-chain peptide can be charged at appropriate pH values, which is not the case with the cyclic peptide. This can provide a basis for separation by electrophoresis.

40. Carbohydrates are not a source of the nitrogen needed for biosynthesis of amino acids.

41. Suggest that your friend show the carboxyl group as a charged carboxylate ($-\text{COO}^-$) and the amino group in its charged form ($-\text{NH}_3^+$).

42. There are very few side chains that have functional groups to form crosslinks.

43. There will be many more possible conformations because of free rotation around the peptide bond.

44. There would be no possibility of disulfide crosslinks within or between peptide chains, giving more possible conformations. There would not be the possibility of oxidation-reduction reactions involving sulfhydryl and disulfide groups.

45. The big difference would be the loss of stereospecificity in the conformation of any peptide or protein. This would have drastic consequences for the kinds of reactions of the protein.

3.5 Are Small Peptides Physiologically Active?

46. Oxytocin has an isoleucine at position 3 and a leucine at position 8; it stimulates smooth muscle contraction in the uterus during

labor and in the mammary glands during lactation. Vasopressin has a phenylalanine at position 3 and an arginine at position 8; it stimulates resorption of water by the kidneys, thus raising blood pressure.

47. The reduced form of glutathione consists of three amino acids with a sulfhydryl group; the oxidized form consists of six amino acids and can be considered the result of linking two molecules of reduced glutathione by a disulfide bridge.

48. Enkephalins are pentapeptides (Y-G-G-F-L, leucine enkephalin, and Y-G-G-F-M, methionine enkephalin), which are naturally occurring analgesics.

49. In most cases, D-amino acids are toxic. They occur in nature in antibiotics and bacterial cell walls.

50. The different stereochemistry of the two peptides leads to different binding with taste receptors and to the sweet taste for one and the bitter taste for the other.

Chapter 4 Answers

4.1 How Does the Structure of Proteins Determine Their Function?

1. (a)(2); (b)(4); (c)(1); (d)(3).

2. When a protein is denatured, the interactions that determine secondary, tertiary, and any quaternary structures are overcome by the presence of the denaturing agent. Only the primary structure remains intact.

3. The random portions of a protein do not contain structural motifs that are repeated within the protein, such as α -helix or β -pleated sheet, but three-dimensional features of these parts of the protein are repeated from one molecule to another. Thus, the term "random" is somewhat of a misnomer.

4.2 What Is the Primary Structure of Proteins?

4. When a protein is covalently modified, its primary structure is changed. The primary structure determines the final three-dimensional structure of the protein. The modification disrupts the folding process.

5. (a) Serine has a small side chain that can fit in any relatively polar environment. (b) Tryptophan has the largest side chain of any of the common amino acids, and it tends to require a nonpolar environment. (c) Lysine and arginine are both basic amino acids; exchanging one for the other would not affect the side chain pKa, in a significant way. Similar reasoning applies to the substitution of a nonpolar isoleucine for a nonpolar leucine.

6. Glycine is frequently a conserved residue because its side chain is so small as to fit in spaces that will not accommodate larger ones.

7. When alanine is replaced by isoleucine, there is not enough room in the native conformation for the larger side chain of the isoleucine. Consequently, there is a great enough change in the conformation of the protein that it loses activity. When glycine is substituted in turn for isoleucine, the presence of the smaller side chain leads to resumption of the active conformation.

8. Meat consists largely of animal proteins and fat. The temperatures

- involved in cooking meat are usually more than enough to denature the protein part of the meat.
- Prion diseases have been linked to the immune system. It is believed that the prion proteins travel in the lymph system bound to lymphocytes and eventually arrive at the nervous tissue where they begin to transform the normal prion protein into the abnormal one.
 - While there are strong genetic predispositions to acquire scrapie, that alone will not cause the disease. The disease must be started by ingesting a prion that already has the altered conformation, PrP^{sc}.
 - The protein efficiency ratio is an arbitrary measurement of the essential amino acid content of a given type of protein.
 - Eggs have the highest PER.
 - The amino acids that must be consumed in the diet because the body cannot synthesize them in sufficient quantities.
 - Reasons for creating genetically modified foods include increasing their protein content, increasing their shelf life, increasing their resistance to insects or other pests, and decreasing the need for pesticides to grow them.

4.3 What Is the Secondary Structure of Proteins?

- Shape, solubility, and type of biological function (static, structural versus dynamic, catalytic).
- The angles of the amide planes as they rotate about the α -carbon. The angles are both defined as zero when the two planes would be overlapping such that the carbonyl group of one contacts the N-H of the other.
- A β -bulge is a common nonrepetitive irregularity found in antiparallel β -sheets. A misalignment occurs between strands of the β -sheet causing one side to bow outward.
- A reverse turn is a region of a polypeptide where the direction changes by about 180° . There are two kinds—those that contain proline and those that do not. See Figure 4.6 for examples.
- The α -helix is not fully extended, and its hydrogen bonds are parallel to the protein fiber. The β -pleated sheet structure is almost fully extended, and its hydrogen bonds are perpendicular to the protein fiber.
- The α -unit, the β -unit, the β -meander, the Greek key, the β -barrel.
- The geometry of the proline residue is such that it does not fit into the α -helix, but it does fit exactly for a reverse turn. See Figure 4.10(c).
- Glycine is the only residue small enough to fit at crucial points in the collagen triple helix.
- The principal component of wool is the protein keratin, which is a classic example of α -helical structure. The principal component of silk is the protein fibroin, which is a classic example of β -pleated sheet structure. The statement is somewhat of an oversimplification, but it is fundamentally valid.
- Wool, which consists largely of the protein keratin, shrinks because of its α -helical conformation. It can stretch and then shrink. Silk consists largely of the protein fibroin, which has the fully extended β -sheet conformation, with far less tendency to stretch or shrink.

4.4 What Can We Say About the Thermodynamics of Protein Folding?

- (1) Backbone H-bonds, involving the CO and NH groups of the peptide chain; (2) side chain H-bonds, involving any possible hydrogen bond donor or acceptors on the side chains; (3) hydrophobic interactions, involving the nonpolar groups on the protein; (4) electrostatic interactions, involving any charged groups on the protein; (5) metal ligation, involving coordination bonds between side chains and a metal ion.
- Stabilization of any conformation depends on lowering of energy, regardless of the kind of interaction involved. With small molecule interactions, the main component of energy changes is enthalpic, with relatively small entropy changes. In the case of molecules as large as proteins, the number of possible conformations makes the entropic part of energy changes much more important. This is particularly true in the case of hydrophobic interactions.

4.5 What Is the Tertiary Structure of Proteins?

- See Figure 4.2 for a hydrogen bond that is part of the α helix (secondary structure). See Figure 4.13 for a hydrogen bond that is part of tertiary structure (side chain hydrogen bonding).
- See Figure 4.13 for electrostatic interactions, such as might be seen between the side chains of lysine and aspartate.
- See Figure 4.13 for an example of a disulfide bond.
- See Figure 4.13 for an example of hydrophobic bonds.
- A chaperone is a protein that aids another protein in folding correctly and keeps it from associating with other proteins before it has reached its final, mature form.
- Configuration refers to the position of groups due to covalent bonding. Examples include *cis* and *trans* isomers and optical isomers. Conformation refers to the positioning of groups in space due to rotation around single bonds. An example is the difference between the eclipsed and staggered conformations of ethane.
- Five possible features limit possible protein configurations and conformations. (1) Although any one of 20 amino acids is possible at each position, only one is used, as dictated by the gene that codes for that protein. (2) Either a D- or an L-amino acid could be used at each position (except for glycine), but only L-amino acids are used. (3) The peptide group is planar so that only *cis* and *trans* arrangements are observed. The *trans* form is more stable and is the one usually found in proteins. (4) The angles ϕ and ψ can theoretically take on any value from 0° to 360° , but some angles are not possible because of steric hindrance; angles that are sterically allowed may not have stabilizing interactions, such as those in the α -helix. (5) The primary structure determines an optimum tertiary structure, according to the “second half of the genetic code.”
- Technically, collagen has quaternary structure because it has multiple polypeptide chains. However, most discussions of quaternary structure involve subunits of globular proteins, not fibrous ones like collagen. Many scientists consider the collagen triple helix to be an example of a secondary structure.

4.6 Can We Predict Folding from Sequence?

- This level of sequence homology is marginal for use of comparative modeling. It is best to try that method, but then to compare the results with those obtained from the fold recognition approach.
- See the Protein Data Bank.
- A prion is a potentially infectious protein found in multiple forms in mammals, often concentrated in nervous tissue. There is a normal form and an abnormal form that tends to form plaques destroying the nervous tissue. They have been found to be transmissible across species.
- A series of encephalopathies have been found to be caused by the abnormal prion protein. In cows, the disease caused by prions is called bovine spongiform encephalopathy or more commonly mad cow disease. In sheep the disease is called scrapie. In humans it is called Jakob-Creutzfeld disease.
- The normal form of the prion protein has a higher α -helix content compared to β -sheet. The abnormal one has an increased β -sheet content.

4.7 What Is the Quaternary Structure of Proteins?

- Similarities—both contain heme group; both are oxygen binding; secondary structure is primarily α -helix. Differences—hemoglobin is a tetramer, while myoglobin is a monomer; oxygen binding to hemoglobin is cooperative, but noncooperative to myoglobin.
- The crucial residues are histidines in both proteins.
- Myoglobin's highest level of organization is tertiary. Hemoglobin's is quaternary.
- The function of hemoglobin is oxygen transport; its sigmoidal binding curve reflects the fact that it can bind easily to oxygen at comparatively high pressures and release oxygen at lower pressures. The function of myoglobin is oxygen storage; as a result, it is easily saturated with oxygen at low pressures, as shown by its hyperbolic binding curve.
- In the presence of H^+ and CO_2 , both of which bind to hemoglobin, the oxygenbinding capacity of hemoglobin decreases.
- In the absence of 2,3-bisphosphoglycerate, the binding of oxygen by hemoglobin resembles that of myoglobin, characterized by lack of cooperativity. 2,3-Bisphosphoglycerate binds at the center of the hemoglobin molecule, increases cooperativity, stabilizes the deoxy conformation of hemoglobin, and modulates the binding of oxygen so that it can easily be released in the capillaries.
- Fetal hemoglobin binds oxygen more strongly than adult hemoglobin. See Figure 4.25.
- Histidine 143 in a β -chain is replaced by a serine in a γ -chain.
- Deoxygenated hemoglobin is a weaker acid (has a higher pK_a) than oxygenated hemoglobin. In other words, deoxygenated hemoglobin binds more strongly to H^+ than does oxygenated hemoglobin. The binding of H^+ (and of CO_2) to hemoglobin favors the change in quaternary structure to the deoxygenated form of hemoglobin.

- The primary flaw in your friend's reasoning is a reversal of the definition of pH, which is $pH = -\log [H^+]$. If the release or binding of hydrogen ion by hemoglobin were the primary factor in the Bohr effect, the pH changes would be the opposite of those actually observed. The response of hemoglobin to changes in pH is the central point. When the pH increases, the hydrogen ion concentration decreases, and vice versa.
- The change of a histidine to a serine in the γ -chain removes a positively charged amino acid that could have interacted with BPG. Thus there are fewer salt bridges to break, so binding is easier than it is in a β -chain.
- Persons with sickle-cell trait have some abnormal hemoglobin. At high altitudes, there is less oxygen, and the concentration of the deoxy form of the abnormal hemoglobin increases. Less oxygen can be bound, causing the observed breathing difficulties.
- In fetal hemoglobin, the subunit composition is $\alpha_2\gamma_2$ with replacement of the β -chains by the γ -chains. The sickle-cell mutation affects the β -chain, so the fetus homozygous for HbS has normal fetal hemoglobin.
- The relative oxygen affinities allow oxygen to be taken by the fetal cells from the maternal Hb.
- Because people with sickle-cell disease are chronically anemic, some cells with fetal Hb are produced to help overcome the impaired oxygen delivery system.
- The crystalline form changed because oxygen entered under the cover slip, transforming deoxyhemoglobin to oxyhemoglobin.

Chapter 5 Answers

5.1 How Do We Extract Pure Proteins from Cells?

- Blender, Potter-Elvehjem, sonicator
- If you needed to maintain the structural integrity of the subcellular organelles, a Potter-Elvehjem would be better because it is more gentle. The tissue, such as liver, must be soft enough to use with this device.
- Salting out is a process whereby a highly ionic salt is used to reduce the solubility of a protein until it comes out of solution and can be centrifuged. The salt forms ion-dipole bonds with the water in the solution, which leaves less water available to hydrate the protein. Nonpolar side chains begin to interact between protein molecules, and they become insoluble.
- Their amino acid content and arrangements make some proteins more soluble than others. A protein with more highly polar amino acids on the surface will be more soluble than one with more hydrophobic ones on the surface.
- Homogenize the liver cells using a Potter-Elvehjem homogenizer. Spin the homogenate at 500 X g to sediment the unbroken cells and nuclei. Centrifuge the supernatant at 15,000 X g, and collect the pellet, which contains the mitochondria.
- No, peroxisomes and mitochondria have overlapping sedimentation characteristics. Other techniques, such as sucrose gradient centrifugation, would have to be used to separate the two organelles.

- If the protein were cytosolic, once the cells were broken open, you could centrifuge at 100,000 X g, and all the organelles would be in the pellet. Your enzyme would be in the supernatant, along with all the other cytosolic ones.
- Isolate the mitochondria via differential or sucrose gradient centrifugation. Use another homogenization technique combined with a strong detergent to release the enzyme from the membrane.
- Tables exist to tell you how many grams of ammonium sulfate (AS) to add to get a certain percent saturation. A good plan would be to take the homogenate and add enough AS to yield a 20% saturated solution. Let the sample sit for 15 minutes on ice and then centrifuge. Separate the supernatant from the precipitate. Assay both for the protein you are working with. Add more AS to the supernatant to arrive at a 40% saturated solution, and repeat the process. In this way, you will find out what the percent saturation in AS needs to be to precipitate it.
- Reasonably harsh homogenization would be able to liberate the soluble protein X from the peroxisomes, which are fragile. Centrifugation at 15,000 X g would sediment the mitochondria (broken or intact). The supernatant would then have protein X but no protein Y. Freeze thaw techniques and sonication would accomplish the same thing, or the mitochondria and the peroxisomes could be separated initially by sucrose gradient centrifugation.

5.2 What Is Column Chromatography?

- (a) Size, (b) specific ligand binding ability, (c) net charge
- The largest proteins elute first; the smallest elute last. Larger proteins are excluded from the interior of the gel bead so they have less available column space to travel. Essentially, they travel a shorter distance and elute first.
- A compound can be eluted by raising the salt concentration or by adding a mobile ligand that has a higher affinity for the bound protein than the stationary resin ligand does. Salt is cheaper but less specific. A specific ligand may be more specific but is often expensive.
- A compound can be eluted by raising the salt concentration or by changing the pH. Salt is cheap, but it might not be as specific for a particular protein. Changing the pH may be more specific for a tight pI range, but extremes of pH may also denature the protein.
- Raising the salt concentration is relatively safe. Most proteins will elute this way and still be active in the case of an enzyme. The salt can be removed later via dialysis if necessary. Changing the pH enough to remove the charge can cause the proteins to become denatured. Many proteins are not soluble at the isoelectric points.
- The basis of most resins is agarose, cellulose, dextran, or polyacrylamide.
- See Figure 5.7.
- Within the fractionation range of a gel filtration column, molecules will elute with a linear relationship of log MW versus their elution volumes. A series of standards can be run to standardize the column, and then an unknown can be determined by measuring its elution volume and comparing to a standard curve.
- Both proteins would elute in the void volume together and would

not be separated.

- Yes, the β -amylase would come out in the void volume, but the BSA would be included in the column bead and would elute more slowly.
- Set up an anion exchange column, such as Q-Sepharose (quaternary amine). Run the column at pH 8.5, a pH at which the protein X has a net negative charge. Put a homogenate containing protein X on the column and wash with the starting buffer. Protein X will bind to the column. Then elute by running a salt gradient.
- Use a cation exchange column, like CM-Sepharose, and run it at pH 6. Protein X will have a positive charge and will stick to the column.
- With a quaternary amine, the column resin always has a net positive charge, and you don't have to worry about the pH of your buffer altering the form of the column. With a tertiary amine, there is a dissociable hydrogen, and the resin may be positive or neutrally charged depending on the buffer pH.
- The easiest way would be to use a sucrose gradient to separate the mitochondria from the peroxisomes first. Then break open the mitochondria via harsh homogenization or sonication. Centrifuge the mitochondria. The pellet would contain protein B, while the supernatant contained protein A. Contaminants could still exist, but they can be cleaned away by running gel filtration, on Sephadex G-75, which would separate C from A and B, and then ion exchange chromatography on Q-Sepharose at pH 7.5. Protein B would be neutral and would elute, while protein A would stick to the column.
- Glutamic acid will be eluted first because the column pH is close to its pI. Leucine and lysine will be positively charged and will stick to the column. To elute leucine, raise the pH to around 6. To elute lysine, raise the pH to around 11.
- A nonpolar mobile solvent will move the nonpolar amino acids fastest, so phenylalanine will be the first to elute, followed by glycine and then glutamic acid.
- The nonpolar amino acids will stick the most to the stationary phase, so glutamic acid will move the fastest, followed by glycine and then phenylalanine.
- A protein solution from an ammonium sulfate preparation is passed over a gel filtration column where the proteins of interest will elute in the void volume. The salt, being very small, will move through the column slowly. In this way, the proteins will leave the salt behind and exit the column without it.

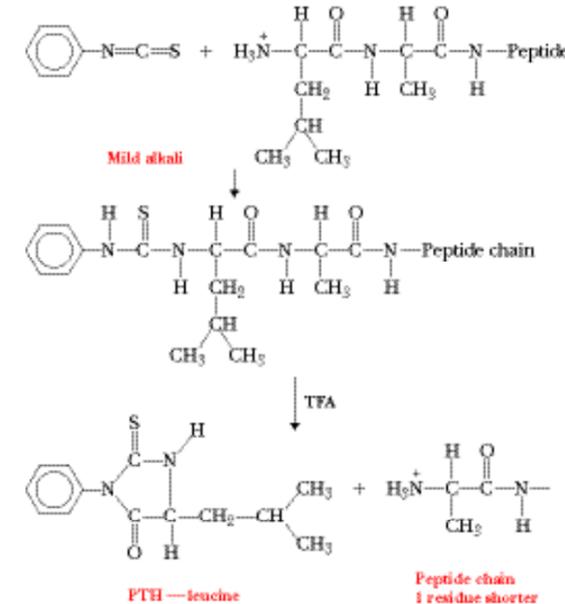
5.3 What Is Electrophoresis?

- Size, shape, and charge.
- Agarose and polyacrylamide.
- Polyacrylamide.
- DNA is the molecule most often separated on agarose electrophoresis, although proteins can also be separated.
- Those with the highest charge/mass ratio would move the fastest. There are three variables to consider, and most electrophoreses are done in a way to eliminate two of the variables so that the separation is by size or by charge, but not by both.

- Sodium dodecyl sulfate polyacrylamide gel electrophoresis. With SDS-PAGE, the charge and shape differences of proteins are eliminated so that the only parameter determining the migration is the size of the protein.
- SDS binds to the protein in a constant ratio of 1.4 g SDS per gram of protein. It coats the protein with negative charges and puts it into a random coil shape. Thus, charge and shape are eliminated.
- In a polyacrylamide gel used for gel filtration chromatography, the larger proteins can travel around the beads, thereby having a shorter path to travel and eluting faster. With electrophoresis, the proteins are forced to go through the matrix, so they travel more slowly if they are bigger because there is more friction.
- The MW is 37,000 daltons.

5.4 How Do We Determine the Primary Structure of a Protein?

- The Edman degradation will give the identity of the N-terminus in its first cycle, so doing a separate experiment is not necessary.
- It might tell you if the protein was pure or if there were subunits.
-



- The amount of Edman reagent must exactly match the amount of N-termini in the first reaction. If there is too little Edman reagent, some of the N-termini will not react. If there is too much, some of the second amino acid will react. In either case, there will be a small amount of contaminating PTH derivatives. This error grows with the number of cycles run until the point that two amino acids are released in equal amounts, and you could not tell which one was supposed to be the correct one.
- In the first cycle, the first and second amino acids from the N-terminal end would be reacted and released as PTH derivatives. You would get a double signal and not know which one was the true N-terminus.
- Val-Leu-Gly-Met-Ser-Arg-Asn-Thr-Trp-Met-Ile-Lys-Gly-Tyr-Met-

Gln-Phe

- Met-Val-Ser-Thr-Lys-Leu-Phe-Asn-Glu-Ser-Arg-Val-Ile-Trp-Thr-Leu-Met-Ile
- It is possible that your protein is not pure and needs additional purification steps to arrive at a single polypeptide. It is also possible that the protein has subunits, so multiple polypeptide chains could be yielding the contradictory results.
- There are two fragments that have C-termini that are not lysine or arginine, which is what trypsin is specific for. Normally there would be only one fragment ending with an amino acid that was not Arg or Lys, and we would immediately know that it was the C-terminus. Histidine is a basic protein, although it is usually neutral and therefore does not react with trypsin. It is possible that in the pH environment of the reaction, the histidine was positively charged and was recognized by trypsin.
- It would tell you a relative concentration of the various amino acids. This is important because it would help you plan your sequencing experiment better. For example, if you had a protein whose composition showed no aromatic amino acids, it would be a waste of time to use a chymotrypsin digestion.
- Cyanogen bromide would be useless as there is no methionine. Trypsin would be little better as the protein is 35% basic residues. Trypsin would shred the protein into over 30 pieces, which would be very hard to analyze.
- Chymotrypsin would be a good choice. There are more than four residues of aromatic amino acids. The 100 amino acid protein would be cut four times, possibly yielding nice fragments in the 20 - 30 amino acid length, which can be sequenced effectively by the Edman degradation.
- It would work best if the basic residues were spread out in the protein. In that way, fragments in the proper size range would be generated. If all four of the basic residues were in the first ten amino acids, there would be one long fragment that could not be sequenced.

Chapter 6 Answers

6.1 What Makes Enzymes Such Effective Biological Catalysts?

- Enzymes are many orders of magnitude more effective as catalysts than are nonenzymatic catalysts.
- The majority of enzymes are proteins, but some catalytic RNAs (ribozymes) are known.
- About 3 seconds (1 year X 1 event/10⁷ events X 365 days/year X 24 hours/day X 3600 seconds/hour = 3.15 seconds).
- Enzymes hold the substrates in favorable spatial positions, and they bind effectively to the transition state to stabilize it. Note that all catalysts lower the activation energy, so this is not a particular enzyme function.

6.2 What Is the Difference between the Kinetic and Thermodynamic Aspects of Reactions?

- The reaction of glucose with oxygen is thermodynamically favored,

as shown by the negative free energy change. The fact that glucose can be maintained in an oxygen atmosphere is a reflection of the kinetic aspects of the reaction, requiring overcoming an activation-energy barrier.

6. First question, most probably; local concentrations (mass action concepts) could easily dictate the direction. Second question, probably not; local concentrations would seldom be sufficient to overcome a relatively large ΔG° of -5.3 kcal in the reverse reaction. (See, however, the aldolase reaction in glycolysis.)
7. Heating a protein denatures it. Enzymatic activity depends on the correct three-dimensional structure of the protein. The presence of bound substrate can make the protein harder to denature.
8. The results do not prove that the mechanism is correct because results from different experiments could contradict the proposed mechanism. In that case, the mechanism would have to be modified to accommodate the new experimental results.
9. The presence of a catalyst affects the rate of a reaction. The standard free energy change is a thermodynamic property that does not depend on the reaction rate. Consequently, the presence of the catalyst has no effect.
10. The presence of a catalyst lowers the activation energy of a reaction.
11. Enzymes, like all catalysts, increase the rate of the forward and reverse reaction to the same extent.
12. The amount of product obtained in a reaction depends on the equilibrium constant. A catalyst does not affect that.

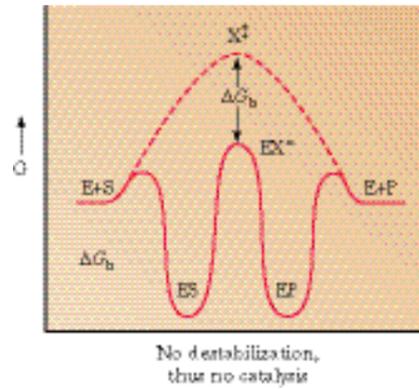
6.3 How Can We Describe Enzyme Kinetics in Mathematical Terms?

13. The reaction is first order with respect to A, first order with respect to B, and second order overall. The detailed mechanism of the reaction is likely to involve one molecule each of A and B.
14. The easiest way to follow the rate of this reaction is to monitor the decrease in absorbance at 340 nm, reflecting the disappearance of NADH.
15. The use of a pH meter would not be a good way to monitor the rate of the reaction. You are probably running this reaction in a buffer solution to keep the pH relatively constant. If you are not running the reaction in a buffer solution, you run the risk of acid denaturation of the enzyme.
16. Enzymes tend to have fairly sharp pH optimum values. It is necessary to ensure that the pH of the reaction mixture stays at the optimum value. This is especially true for reactions that require or produce hydrogen ions.

6.4 How Do Substrates Bind to Enzymes?

17. In the lock-and-key model, the substrate fits into a comparatively rigid protein that has an active site with a well-defined shape. In the induced-fit model, the enzyme undergoes a conformational change on binding to substrate. The active site takes shape around the substrate.

18.



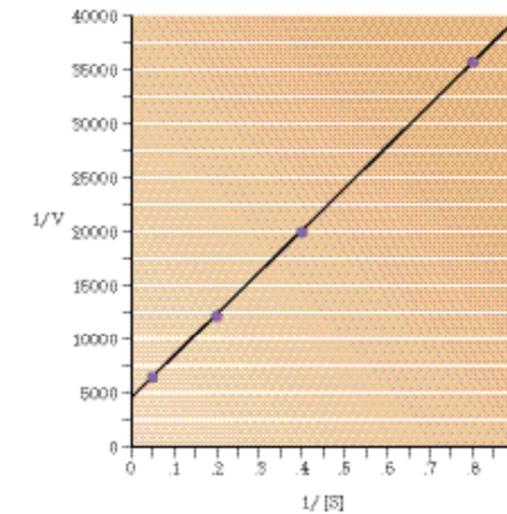
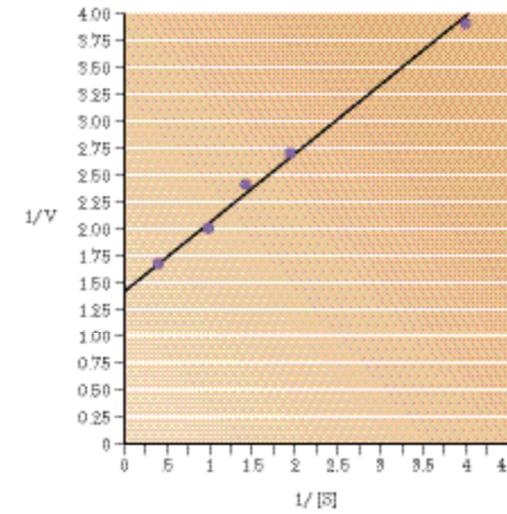
19. The ES complex would be in an "energy trough," with a consequentially large activation energy to the transition state.
20. Amino acids that are far apart in the amino acid sequence can be close to each other in three dimensions because of protein folding. The critical amino acids are in the active site.
21. The overall protein structure is needed to ensure the correct arrangement of amino acids in the active site.
22. The strong inhibition indicates tight binding to the active site. Thus, the compound is very likely to be a transition state analogue.

6.5 What Are Some Examples of Enzyme-Catalyzed Reactions?

23. See Figures 6.6 and 6.7.
24. Not all enzymes follow Michaelis-Menten kinetics. The kinetic behavior of allosteric enzymes does not obey the Michaelis-Menten equation.
25. The graph of rate against substrate concentration is sigmoidal for an allosteric enzyme but hyperbolic for an enzyme that obeys the Michaelis-Menten equation.

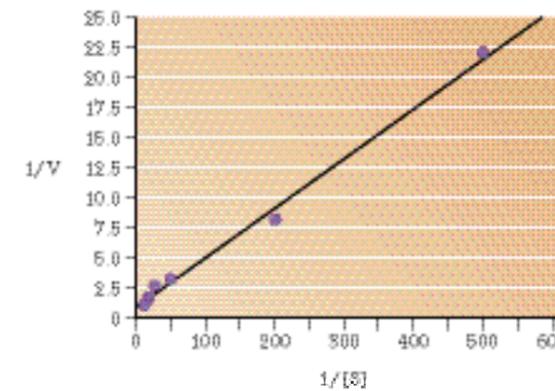
6.6 What Is the Michaelis-Menten Approach to Enzyme Kinetics?

26. The reaction velocity remains the same with increasing enzyme concentration. It is theoretically possible, but highly unlikely, for a reaction to be saturated with enzyme.
27. The steady-state assumption is that the concentration of the enzyme-substrate complex does not change appreciably over the time in which the experiment takes place. The rate of appearance of the complex is set equal to its rate of disappearance, simplifying the equations for enzyme kinetics.
28. Turnover number = $V_{max}/[E]_t$.
29. Use Equation 6.16, (a) $V = 0.5 V_{max}$; (b) $V = 0.33 V_{max}$; (c) $V = 0.09 V_{max}$; (d) $V = 0.67 V_{max}$; (e) $V = 0.91 V_{max}$.
30. See graph: $V_{max} = 0.681 \text{ mMmin}^{-1}$, $K_M = 0.421 \text{ M}$.

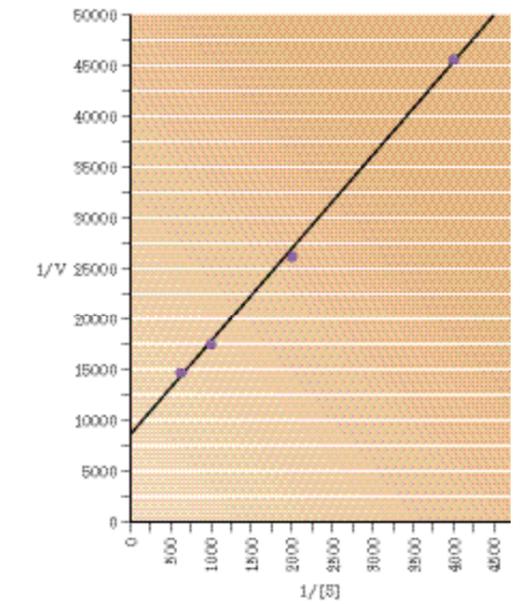


31. See graph: $V_{max} = 2.5 \times 10^{-4} \text{ Msec}^{-1}$, $K_M = 1.6 \times 10^8 \text{ M}$.

32. See graph: $K_M = 2.86 \times 10^{-2} \text{ M}$. Concentrations were not determined directly. Absorbance values were used instead as a matter of convenience.



33. See graph: $V_{max} = 1.32 \times 10^{-3} \text{ Mmin}^{-1}$, $K_M = 1.23 \times 10^{-3} \text{ M}$.



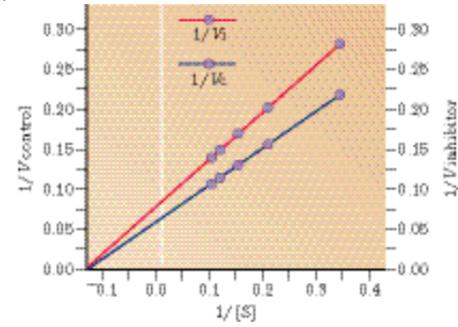
34. The turnover number is 20.43 per minute.
35. The number of moles of enzyme is 1.56×10^{-10} . Turnover number = $10,700 \text{ sec}^{-1}$.
36. The low K_M for the aromatic amino acids indicates that they will be oxidized preferentially.
37. It is easier to detect deviations of individual points from a straight line than from a curve.
38. The assumption that the K_M is an indication of the binding affinity between the substrate and the enzyme is valid when the rate of dissociation of the enzyme-substrate complex to product and enzyme is much smaller than the rate of dissociation of the complex to enzyme and substrate.

6.7 How Do Enzymatic Reactions Respond to Inhibitors?

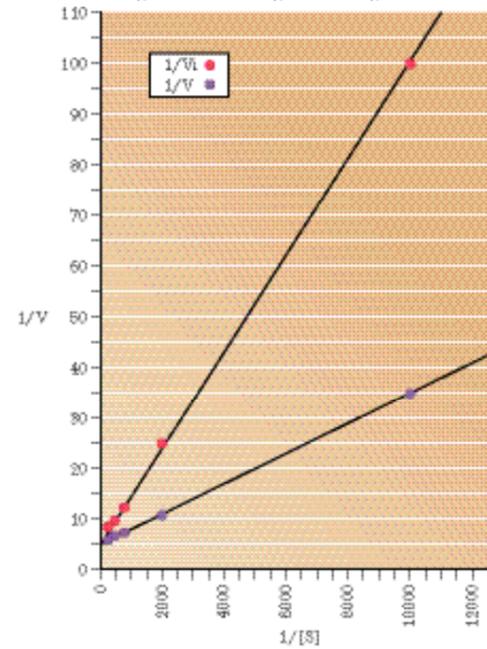
39. In the case of competitive inhibition, the value of K_M increases, while the value of K_M remains unchanged in noncompetitive inhibition.
40. A competitive inhibitor blocks binding, not catalysis.
41. A noncompetitive inhibitor does not change the affinity of the enzyme for its substrate.
42. A competitive inhibitor binds to the active site of an enzyme, preventing binding of the substrate. A noncompetitive inhibitor binds at a site different from the active site, causing a conformational change, which renders the active site less able to bind substrate and convert it to product.
43. Competitive inhibition can be overcome by adding enough substrate, but this is not true for all forms of enzyme inhibition.
44. A Lineweaver-Burk plot is useful because it gives a straight line. It is easier to determine how well points fit to a straight line than to a curve.
45. In a Lineweaver-Burk plot for competitive inhibition, the lines intersect at the y-axis intercept, which is equal to $1/V_{max}$. In a Lineweaver-Burk plot for noncompetitive inhibition, the lines

intersect at the x-axis intercept, which is equal to $-1/K_M$.

46. $K_M = 7.42 \text{ mM}$; $V_{\max} = 15.9 \text{ mmol min}^{-1}$; noncompetitive inhibition.



47. Competitive inhibition, $K_M = 6.5 \times 10^{-4}$. The key point here is that the V_{\max} is the same within the limits of error. Some of the concentrations are given to one significant figure.



48. It is very good in the case of noncompetitive inhibitors; much of metabolic control depends on feedback inhibition by downstream noncompetitive inhibitors. The question is perhaps moot in the case of competitive inhibitors, which are much less commonly encountered *in vivo*. Some antibiotics, however, are competitive inhibitors (good for the sick person, bad for the bacteria).
49. Both the slope and the intercepts will change. The lines will intersect above the x-axis at negative values of $1/[S]$.
50. Not all AIDS drugs are enzyme inhibitors, but an important class of such drugs inhibits the HIV protease. You would need to understand the concepts of substrate binding, inhibition, and inhibitor binding.
51. An irreversible inhibitor is bound by covalent bonds. Noncovalent interactions are relatively weak and easily broken.
52. A noncompetitive inhibitor does not bind to the active site of an enzyme. Its structure need bear no relation to that of the substrate

Chapter 7 Answers

7.1 Does the Michaelis - Menten Model Describe the Behavior of Allosteric Enzymes?

- Allosteric enzymes display sigmoidal kinetics when rates are plotted versus substrate concentration, Michaelis-Menten enzymes exhibit hyperbolic kinetics. Allosteric enzymes usually have multiple subunits, and the binding of substrates or effector molecules to one subunit changes the binding behavior of the other subunits.
- It is an enzyme used in the early stages of cytidine nucleotide synthesis.
- ATP acts as a positive effector of ATCase and CTP acts as an inhibitor.
- The term K_M should be used for enzymes that display Michaelis - Menten kinetics. Thus, it is not used with allosteric enzymes. Technically, competitive and noncompetitive inhibition are also terms that are restricted to Michaelis - Menten enzymes, although the concepts are applicable to any enzyme. An inhibitor that binds to an allosteric enzyme at the same site as the substrate is similar to a classical competitive inhibitor. One that binds at a different site is similar to a noncompetitive inhibitor, but the equations and the graphs characteristic of competitive and noncompetitive inhibition don't work the same way with an allosteric enzyme.
- A K system is an allosteric enzyme in which the binding of inhibitor alters the apparent substrate concentration needed to reach one-half V_{\max} $S_{0.5}$.
- A V system is an allosteric enzyme in which the binding of inhibitor changes the V_{\max} of the enzyme but not the $S_{0.5}$.
- Homotropic effects are allosteric interactions that occur when several identical molecules are bound to a protein. The binding of substrate molecules to different sites on an enzyme, such as the binding of aspartate to ATCase, is an example of a homotropic effect. Heterotropic effects are allosteric interactions that occur when different substances (such as inhibitor and substrate) are bound to the protein. In the ATCase reaction, inhibition by CTP and activation by ATP are both heterotropic effects.
- ATCase is made up of two different types of subunits. One of them is the catalytic subunit, and there are six of them organized into two trimers. The other is the regulatory subunit, which also consists of six protein subunits organized into three dimers.
- Enzymes that exhibit cooperativity do not show hyperbolic curves of rate versus substrate concentration. Their curves are sigmoidal. The level of cooperativity can be seen by the shape of the sigmoidal curve.
- Inhibitors make the shape of the curve more sigmoidal.
- Activators make the shape of the curve less sigmoidal.
- $K_{0.5}$ is the substrate concentration that leads to half of the maximal velocity. This term is used with allosteric enzymes where the term K_M is not appropriate.
- A mercury compound was used to separate the subunits of ATCase. When the subunits were separated, one type of subunit retained catalytic activity but was no longer allosteric and was not inhibited by CTP. The other subunit type had no ATCase activity,

but it did bind to CTP and ATP.

7.2 What Are the Models for the Behavior of Allosteric Enzymes?

- In the concerted model, all the subunits in an allosteric enzyme are found in the same form, either the T or the R. They are in equilibrium with each enzyme having a characteristic ratio of the T/R. In the sequential model, the subunits change individually from T to R.
- The sequential model can explain negative cooperativity because a substrate binding to the T form could induce other subunits to switch to the T form, thereby reducing binding affinity.
- Greater cooperativity is favored by having a higher ratio of the T/R form. It is also favored by having a higher dissociation constant for the substrate binding to the T form.
- The L-value is the equilibrium ratio of the T/R form. The c value is the ratio of the dissociation constants for substrate and the two forms of enzyme, such that $c = KR/KT$.
- Many models are possible. We never really know for sure how the enzyme works, rather we create a model that explains the observed behavior. It is very possible that another model would do so as well.

7.3 How Does Phosphorylation of Specific Residues Regulate Enzyme Activity?

- A kinase is an enzyme that phosphorylates a protein using a high-energy phosphate, such as ATP, as the phosphate donor.
- Serine, threonine, and tyrosine are the three most often phosphorylated amino acids in proteins that are acted upon by kinases. Aspartate is another one often phosphorylated.
- The allosteric effect can be faster because it is based on simple binding equilibrium. For example, if AMP is an allosteric activator of glycogen phosphorylase, the immediate increase in AMP when muscles contract can cause muscle phosphorylase to become more active and provide energy for the contracting muscles. The phosphorylation effect requires the hormone cascade beginning with glucagon or epinephrine. There are many steps before the glycogen phosphorylase is phosphorylated, so the response time is slower. However, the cascade effect produces many more activated phosphorylase molecules, so the effects are longer and stronger.
- As part of the mechanism, the sodium-potassium ATPase has an aspartate residue that becomes phosphorylated. This phosphorylation alters the conformation of the enzyme and causes it to close on one side of the membrane and open on the other, moving ions in the process.
- Glycogen phosphorylase is controlled allosterically by several molecules. In the muscle, AMP is an allosteric activator. In the liver, glucose is an allosteric inhibitor. Glycogen phosphorylase also exists in a phosphorylated form and an unphosphorylated form, with the phosphorylated form being more active.

7.4 What Are Zymogens, and How Do They Control Enzyme Activity?

- The digestive enzymes trypsin and chymotrypsin are classic examples of regulation by zymogens. The blood clotting protein thrombin is another.
- Trypsin, chymotrypsin, and thrombin are all proteases. Trypsin cleaves peptide bonds where there are amino acids with positively charged side chains (Lys and Arg). Chymotrypsin cleaves peptides at amino acids with aromatic side chains. Thrombin cleaves the protein fibrinogen into fibrin.
- The zymogen prothrombin is cleaved to give the active enzyme thrombin. The thrombin then cleaves a soluble molecule, fibrinogen, into an insoluble molecule, fibrin. Fibrin is a protein that forms part of the blood clot.
- Chymotrypsinogen is an inactive zymogen. It is acted upon by trypsin, which cleaves peptides at basic residues, like arginine. When trypsin cleaves between the arginine and the isoleucine, chymotrypsinogen becomes semiactive, forming π -chymotrypsin. This molecule digests itself further, forming the active α -chymotrypsin. As it turns out, the α -amino group of the isoleucine produced by the first cleavage is near the active site of α -chymotrypsin and necessary for its activity.
- Zymogens are often seen with digestive enzymes that are produced in one tissue and used in another. If the enzyme were active immediately upon production, it would digest other cell proteins where it would cause great damage. By having it produced as a zymogen, it can be safely made and then transported to the digestive tissue, such as the stomach or small intestine, where it can then be activated.
- This allows for a more rapid response when the hormone is needed. The hormone is already synthesized and usually just requires breaking one or two bonds to make it active. The hormone can be poised and ready to go on demand.

7.5 How Do Active Site Events of an Enzyme Affect the Reaction Mechanism?

- Serine and histidine are the two most critical amino acids in the active site of chymotrypsin.
- The initial phase releases the first product and involves an acyl-enzyme intermediate. This step is faster than the second part where water comes into the active site and breaks the enzyme-acyl bond.
- In the first step of the reaction the serine hydroxyl is the nucleophile that attacks the substrate peptide bond. In the second step, water is the nucleophile that attacks the acyl-enzyme intermediate.
- His57 performs a series of steps involving general base catalysis followed by general acid catalysis. In the first phase, it takes a hydrogen from Ser195, acting as a general base. This is followed immediately by an acid catalysis step where it gives the hydrogen to the amide group of the peptide bond that is breaking. A similar scheme takes place in the second phase of the reaction.
- The first phase is faster for several reasons. The serine at position

195 is a strong nucleophile for the initial nucleophilic attack. It then forms an acyl enzyme intermediate. In the second phase, water is the nucleophile, and it takes time for water to diffuse to the right spot to perform its nucleophilic attack. It is also not as strong a nucleophile as the serine. Therefore, it takes longer for water to perform its nucleophilic attack and break the acyl-enzyme intermediate than it takes for serine to create it.

- 35. His57 exists in both the protonated and unprotonated form during the chymotrypsin reaction. Its pKa of 6.0 makes this possible in the physiological pH range.
- 36. Instead of a phenylalanine moiety (similar to the usual substrates of chymotrypsin), use a nitrogen-containing basic group similar to the usual substrates of trypsin.

7.6 What Types of Chemical Reactions are Involved in Enzyme Mechanisms?

- 37. They act as Lewis acids (electron pair acceptors) and can take part in enzyme catalysis mechanisms of enzymes.
- 38. False. The mechanisms of enzymatic catalysis are the same as those encountered in organic chemistry, operating in a complex environment.
- 39. General acid catalysis is the part of an enzyme mechanism where an amino acid or other molecule donates a hydrogen ion to another molecule.
- 40. S_N1 stands for unimolecular nucleophilic substitution. The unimolecular part means that it obeys first-order kinetics. If the reaction is R-X + Z: → R-Z + X:, with an S_N1 reaction, the rate is dependent on the speed with which the X breaks away from the R. The Z group comes in later and quickly, compared to the breakdown of R-X. S_N2 stands for bimolecular nucleophilic substitution. This happens with the same reaction scheme if the Z attacks the R-X molecule before it breaks down. Thus the concentration of both R-X and Z: are important, and the rate displays second-order kinetics.
- 41. The S_N1 reaction leads to loss of stereospecificity as the X group leaves before the entering nucleophile. This means that the nucleophile can enter from different angles leading to different isomers.
- 42. The results do not prove that the mechanism is correct because results from different experiments could contradict the proposed mechanism. In that case, the mechanism would have to be modified to accommodate the new experimental results.

7.7 What Is the Connection between the Active Site and Transition States?

- 43. A good transition state analogue would have to have a tetrahedral carbon atom where the amide carbonyl group was originally found, since the transition state involves a momentary tetrahedral form. It would also have to have oxygens on the same carbon so that there would be sufficient specificity for the active site.
- 44. The induced fit model assumes that the enzyme and substrate must both move and change to conform to each other perfectly. Thus, the true fit is not between the enzyme and substrate but

between the enzyme and the transition state of the substrate on its way to product. A transition state analogue will fit the enzyme nicely in this model.

SW. Rs rOPROB LE MOCHAPTEr 1A|swelsj | I wa take th eBS i chemisfor thistextij AP ol ymerisive r' laræmol. CUi Eror meyl i li rnsM auifufits (moflo llers)to ætherA, rotinis afolynefro rfid byi nçinlla NaCias toeth efi A nlc le ic acidi s apoymerfor m eaby- iling nicle. ti a estogæte le n cataly si ssthePr oës stiatijr

ASstheritel fchemiclar Eaction SconPa rca tothpulle atily zeaeaction ftoio gicau lat at ystarep. Oteia Signaibos auic is (S: thonu. Vexcep, ti onstleafeityp esO ERNA>w na C hcaflaT ai yze someof terele it ion Sorth eio lnnct a fOlislm. Selet i Coed ei Stnell e ansbyth chde in f0 rm au offo rnestr uCturalna full Cti0n ofra lily jnst hifng sis pas ed f rjmoles ener a ti o

nto. heneiu me Se QuencOf Pu r ipesllp yrjpi a ifles inDvalgri es tere Neti (Coile rma st recodi n' materal in n, Om eVi puses) u l 2Wha t jSstherim ai naufo erip jra nbiom0 i tcaud) r tcor rct rdtcpof lin ction. Isopolan acPolukas colita ininistat utcti ofiAs fO lbi sga le ni ntefoi loWnsjis u amino srOU P CHC:Hz 2 Cq rOn- isro p rcketon-) CH3C0: HBY ar0 XYsr0UP C:H0Hj amox. I gto U rC:H3Co0i (Ar fonyler oupc al d'hyd eCH3Hz2 CHOpi0L. rO p rC: 3SHjSTer i il idg eCH3COO: HCH3lou hæfou, CH3 (HC HCH)Aidæ iikæe CH:Co NCH3>2Hth erCH3H2..

C,2 CH3I the funci ord
18. Ollrsitue (om fOunds f Ollow a ilfo, ei Rstie-onCe r0rvj thi sm hEaitha tWraali c0mP0 undSolia-niYbællæb yia vi rgs Ys tenn anawefese yondtref qAmoflato, atorv inlestisat i0nslw, hie f ss' n hEsi s j0wæ uat, ræll icomfo..

dS. Iiæ i norgnic one Sraimtreq ui reayitali st icll plonæti. N> utra tte, tæy0veæth elawof Ciemst ræna Ph ysiCsand thus rEr e s hæct t0iæto, atorv i nVestis æli on s uæCq uen u y th
e
CO. C0rwa s EX te nde atone ml C p0 fccol pley buts, i lte Stæie la isci pæ fcc0 hio chemisry.

51. ræw i kæa no ræni C0mP 0ln as> h a sthesæm o l e c Ula rstru ctillæ Wh tæ h eræti sPr. duæ eaby qli, i n s0 f' an s n0 u 6. 7 i f- ves7 i rth et Wocy. lo pto rð n eæriya t, vesæ re æll owed s t h i r- Cendæ r' rEta k: o j0is> I i qidæy, es/ ket on e s> æll oællæ ep oxidæsan æt hærslæ.

hæ. Can, i0æe mistf Ys a yæb0rpos. bteoris ilis oæiæp> 9 i t i s g ælæa uYiæiv edth æt carboll isth eliæ reiyææ, is f O rællæ fer omis> æart h0o unæio fæxtæterf æstræ æll i O E i s itæc ræf ælles. Ollæ sive2, i0r 2j 6: i læp ossibi i i t i es' t h u s i r.

Si æw 0 Uææene

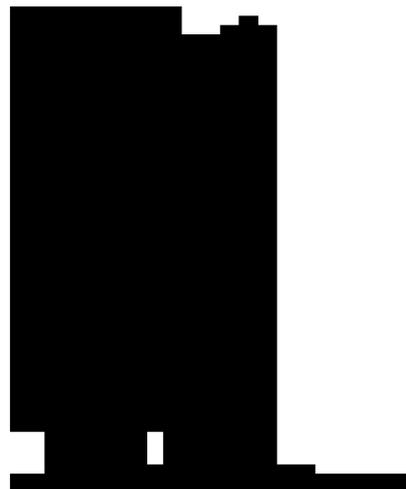
C,SS æ r' to iæv ætæstæv o s a d r o S n
U. h æ r' (æ 2 x iæ 3) o r p l e s s i i i i e s i i i t h e n u m h e r i s æ o r i t h 2 x i o 2 æ h æ- C h i s t w i c e æ l o s æ d o r s i n u l l. æ r i 1 æ r æ i s c æ p æ b æ o f b æ t i c o d i n æ r æ d
C

8.2 What Are the Chemical Natures of the Lipid Types?

- 2. In both types of lipids, glycerol is esterified to carboxylic acids, with three such ester linkages formed in triacylglycerols and two in phosphatidyl ethanolamines. The structural difference comes in the nature of the third ester linkage to glycerol. In phosphatidyl ethanolamines, the third hydroxyl group of glycerol is esterified not to a carboxylic acid, but to phosphoric acid. The phosphoric acid moiety is esterified in turn to ethanolamine. (See Figures 8.2 and 8.5.)
- 3.

- 4. Both sphingomyelins and phosphatidylcholines contain phosphoric acid esterified to an amino alcohol, which must be choline in the case of a phosphatidyl choline and may be choline in the case of a sphingomyelin. They differ in the second alcohol to which phosphoric acid is esterified. In phosphatidylcholines, the second alcohol is glycerol, which has also formed ester bonds to two carboxylic acids. In sphingomyelins, the second alcohol is another amino alcohol, sphingosine, which has formed an amide bond to a fatty acid. (See Figure 8.6.)
- 5. This lipid is a ceramide, which is one kind of sphingolipid.
- 6. Sphingolipids contain amide bonds, as do proteins. Both can have hydrophobic and hydrophilic parts, and both can occur in cell membranes, but their functions are different.
- 7. Any combination of fatty acids is possible.

- 8. Steroids contain a characteristic fused-ring structure, which other lipids do not.
- 9. Waxes are esters of long-chain carboxylic acids and long-chain alcohols. They tend to be found as protective coatings.
- 10. Phospholipids are more hydrophilic than cholesterol. The phosphate group is charged, and the attached alcohol is charged or polar. These groups interact readily with water. Cholesterol has only a single polar group, an -OH.
- 11.



- 12. The waxy surface coating is a barrier that prevents loss of water.
- 13. The surface wax keeps produce fresh by preventing loss of water.
- 14. Cholesterol is not very water soluble, but lecithin is a good natural detergent, which is actually part of lipoproteins that transport the less soluble fats through the blood.
- 15. The lecithin in the egg yolks serves as an emulsifying agent by forming closed vesicles. The lipids in the butter (frequently triacylglycerols) are retained in the vesicles and do not form a separate phase.
- 16. The removal of the oil also removes the natural oils and waxes on the feathers. They must regenerate before the birds can be released.

8.3 What Is the Nature of Biological Membranes?

- 17. Triacylglycerols are not found in animal membranes.
- 18. Statements (c) and (d) are consistent with what is known about membranes. Covalent bonding between lipids and proteins [statement (e)] occurs in some anchoring motifs, but is not widespread otherwise. Proteins “float” in the lipid bilayers rather than being sandwiched between them [statement (a)]. Bulkier molecules tend to be found in the outer lipid layer [statement (b)].
- 19. The public is attuned to the idea of polyunsaturated fats as healthful. The *trans*-configuration gives a more palatable consistency. Recently, however, concerns have arisen about the extent to which such products mimic saturated fats.
- 20. Partially hydrogenated vegetable oils have the desired consistency for many foods, such as oleomargarine and components of TV dinners.
- 21. Many of the double bonds have been saturated. Crisco contains “partially hydrogenated vegetable oils.”
- 22. There is less heart disease associated with diets low in saturated fatty acids.
- 23. The transition temperature is lower in a lipid bilayer with mostly unsaturated fatty acids compared to one with a high percentage of saturated fatty acids. The bilayer with the unsaturated fatty acids is already more disordered than the one with a high percentage of

fe. XPe riefc iil gaSiarrhcia pael c e t h e p l i n t i n a t i t f a t, o n c u r V. Wh. fe u e l l a e d a g d o b a s e e u a u s t l e a m o u n t Δ O ^α u. fe. 0 f i s i n a l l y p r e s e n t (c f) h e p r o b f y o r a p p o i e c u Δ I ^α e. a t s f e a d i n s o u j b i e j n v i t e f i p i e a 4 # s e e i i i i c h i p u r r i s o t i m p s i e s Δ b ^α f e n i n l ^α . a c t y f h a . t i e m i t . j l u j e b u e f e r s h a v e t i e Δ S ^α u . H j i n e d f y a u n a v e t h e s i m e r e i a j v e a p p o u n t s Δ O ^α e . e a c d a n a i s f e f o m 4 l . t w o u j d e m o r e f e e c t y e t o s t a r t Δ i ^α e . e . e p s i s ^α e i . o u w a n t q u i f f e r a f a p h a v o l t e r p a w h i c h i l l e a n s h a t t h e b a s e f f o m w a l i p r e d o m i n a t e w h e n y o u a v e r i n g s h e d p l e n i n . i t i s a s i e r t o c o l l e v e t s o m e o r u e t a s e f o r m o t h e a i c a r o i n t h a m o s . o f . h e a g d o m p l o t h e b a s e e o m p l a o i n a b u f Δ f ^α w . t t h e p h a b o y e e h e p k a t Δ h ^α a . e f . f i p f e a p p i m a t s i t h i s o u u l a p u s e f u l l s Δ a ^α e . e r . o f i j r

ef r e s o r t h e l a v i a u d i a m i n o l - i a s ? 2 i p r o t i n i s t e c h n i c a l l y n o t m a p p i n c a c i . j g i v c a f e . o n t a i p s n o c h . f . i c . t h o s a u i s i s i a n a m i f o a G i d i n v a i c h e r e k e r l l p c o n t a i n s t h e r o i i o w i n g : a n d o v l l g r o u p c e f i n e t h e . n i n e v t y r o s i l l e a s u p p u r a t o m e c y s t e i e r y m e t h o i n e s a s e c o n d e m a i c l f b o n a t o m e i s o , e l c i n b t h r e o g n i e d a m a l i o g r o u p c o u s a r e l a n . m i . e g r o u p a s p a f a f i n e s g u t q u i n e b a l l a c i a s f o u p e d s f a r e a t e s t u e a t e d m a r o m a t e r i n g c o n e p y , q u a l i t y t y r o s i n e t t f . p r o p i a p p a i r a n c h e a s i d e c h a i n k l e u c i n e v a l i n e > i n t e r .

P t d e v i l l e t e s e r i i e n t e l l a r g o s m f r a n e u t e r l a **I** a m . n o a c i d a r i s e n a f s c y s t a n a t y r t h e r o m a t a c a m i n o g a c i d a r . t h e a n t i f a n a t h e s u p f u r l l o l t a j n i s a l l i n a c i a s a r e m e t a n c y s s i i n t e r e p t a e s G u i t h r o m a s h i i n e r e m a i n u e r o n . o l . f a m i l o a c i s a f e v a u i t p a n a i a i t h e a c i d e a i n o a c i d a r e g . u l n a s p i 6 i a m a n a c i a s t h e r t h a n n e h s i i z o a r e r a t e a s y l l a i f i c a t i o n r o f o n e r t h e c o m m o d a m i n o a c i d s e e n i s u r t . 1 5 . o f t h e s t r u c t u r e s o - s o m e t h o a i f i e a d a l l i n o a c i d s i h y a p o x y l u i l l a n a l y d o x m y s i n e a r e f o u l l i n c o l l a g e n t h y f o x i n e i s f o u l i n h y r o s i 0 b l l i j n 7 i r t . s i . e t r y p t o p h a n a n a t h e r l e n y a t v . s i s a m l o p o - m i n o x i a a - e i s a n e n z i m e t h a t e g r a d s c o l l i n u p a s w i t h a n i m i n o s r o u p i r e . u i n s e r l l o r t a n s m i t e r s : c o n s e u e n t i y t h e g y d s c o l l e r o u p e r s . i s m e n q i s t a t e s i t h e n i s h o c o n c e n t r a t i o n o f t r y p t o p h a n i n k e f o u t i n g y m i n d i v e l e v a t e t h e l e v e l s o f s e f o l l i n w h a c h e r a X . s t . e b r a i n i o l t h e f y r p r a l i n m i l k m i s h e n d k e t o u s i e p y w h e f e a s u e t y r i n i n e j n c h e s s h o l l i a r e p o u . i i i t e d m i n a c i a s t h y o x i n e a n i s v a r . x y p r o l . n e o c u f i n v e r y a e w i r o t e i l s i t h e s e n e t c e d

Edoesno ti nciuaet

h e j s o t h e v a f e p r o d u c e s y m o d i f i c a t i o n o f t y r o s  n a p a l p r o i n e r t s p e c t i v e j y i z o s d l a m i n o a c i s . a v e s p e i f i c - c i a . b a s e r o p e r t i e s e i s t a e q l e i m j a z o e i s e d p r o t o n a t e a m . n o s e r o u p i s p r e a m i n a n t i v e r o t o l a t e d i s p a - a s i n e . a m i n o g e t o l i s d e p l o t i n a t e a . f y r o p h a n . i n a n j r o s r o u p i s p l e i o m i t a t e i y a p o t o l a t e d i h r o i i n d i n i n o s t u p i s p a . t i a l l y a e p r o t o n a t e d i t y r o s i l e . a m i n o s r o u p i s e a o m i n a t i y a e p r o t o m t e a p l e n o l i c h y a p o x y l . s a p p r o x i m i t e l y a d s o i s o m i x t u f t h e p r o t o m a t p a l n a d e p r o t o n a . e a i n l j i s o e r e c t i f i c t o i n e r o r s u l t a m i c d c j o z i s s e r a i l e s i z h i s t e j d i n e s 7 5 s t y s i n e s 9 i 7 5 v f o s i l e s i 6 s a f e s i n i . e j l o 7 s j i d c l s t e i n e w a l l h a v e n o n e t c h a f e a d t e p h s l o z e c i i 7 i + 8 3 3 / 2 .

e a c i o n t h a t p l o a l l c e s h t e c a l l s e t t e r e w i l l b e p r i e n t y o u n d a s e r o m t o r e a c t i v i t h e n e h y a r o s n i o p r o l a c e o 4 7 i z w i t t e r i o l s t e . n d m o t o i n t e f f e r e i t h h i o c h e m i c a l i f e a c t i o n s i 4 8 i i t i s u s e r u l t o n a . l a s l i f e r e h a t y i l h a i n t a i n a s t a b l e p e . e n . f a s s a b c o n e i t i o i s c h a n g e . i d i l u t i o n i s o p e s i l t h r o s i s i b e c h a n g e l 4 9 i t a s u s e r u l t o h a v e a b l e f e r t h a t w i m p i n e i n g s t a b l e e n l e v e r i

a s . a v c o n d i t i o n s c h a n g e a t e l l e p e r a u t e v a r i a t i o n i s . n e s u c h p o s s i - b i l e c l a r k i 5 0 i t h t o n i y z w i t t e . i i o n i s + h s t e m m c h a p e r s a n s w - p s o i w h a t a r e m i n o a c i s . a n . w h a t i s t h e l i p h e o d i m e i s i o a m i i m a n d i m a l i n d e i d s h a l e a i r e l r e n e s t e r e d e l e i s t e r y q u o i a c h a n d i n . n l . e p t i e a s t y d . c o n t a i n o m a m i n o a c i a s a r e f o u n d i n b a c t e r i a i c e i t h a i j s a n j r o m e a n t e i b i o t i c s i 3 i 2 w a t f e e s e r u l t i l r e s a n d p r o

e . o b i c o l i a t i o n o f s a u c o s e i s e x e r s i n c a d a s - s o u r c e o f e n e r g y f o r m . n o r a n i s m s i n i u a i n q u i n g s i n w u . a t e r t a s o n - h u e t o e x e r t i e - w o . f o c e s g e s t o t a k e - j l e c a i f r e f e l l . m i l b r a e r t o p r o v i d e p e r s o r t h e e l e k t r o n c o n . i c h a p t e 2 a n s w e r s 2 i h a t s a h a t o g e n e r a t e i i p r o t e i n s a n a n u c l e i c a c i a s h a e n . d . o s e n o n a s a d p i m p o r a n t a r t h e e i r s t r u c t u f e s i r e p u c l i o n o r d n a g n a i s t e j . s e f i p u l l o r n o r n a r e a u i r e s i v a f o s e n o n a i n s o f o m p l e m m a r y h a s e s t o t h e d a t e m p i d t e s t r a n d l y t r e c - h . o . d i s n o t s u f f i c i e n t y r o m a r o l e f e a t y u e t u d i d i s t r i b u t i o n o f e i - c i r o t s a t i s t a y o e l a s t . i s o t h e r a n o u n s h a r e a p a i f s o i e l e c t r o n s t o s e r v e a s y a r e n o n d e c e p t o r s i a m a n y m o l e c u l e s a n f o r m i n a f o l e n i o l i s e x a m p l e s i n i s h t y e l l o c h e s . h . n h a s i r o m a t o t o e c a l i . e t a h y d r o g . n p o n a e i l l u s t r a v e d . v a f o s e i c o r . a u e n e l y b o h e d t o o w .



f . i - h i s h u r t s e n t e n f o n s q i v d f o e n s o n a l i t a m t h e r o n o r t h e a l l a - y a r o s e n . b o l n e a d i m e f o g l e t i c a c i a t h e " o h o r t i o n o r t h e a n d x y i a l l p o n n o i e c - p r a i s i v i . o s e i i s o n a e t t u i e j p o r t i o - o r a h t a l t o x y l s r o u p d i m o l . c u . e a d a t i v e e v e r s q l z i e i l l o s e a . a n d o n i t o i i s l i b i o l i s e a c i a l d o i n g o l l e a b o n a t o s w a t e r a l t h e f i n s o x i e n s i l l a s t o w d t r - s

u g . f a u c o m i s b i n q m r e t h a n t h e c o r r e s p o n d i n s u f a r s l o s h i p o v e t i a i a t i o n e c e a s e s t . e p h o m c o d i c s e e b c o n p 5 a d 9 a s p i n i s e i - t e r i c a l y r e u r a i . t t . e p h o t h e s t o i a c i a n a c i a s s i e m e m p r a n e m o f e e a s i y t r e r e t h a n i l - h s l l a i n t e s t a n e l j o r t o s i t i v e y c h a n g e e a o m s . i l i b i n d i t o n l - i d a c a g s i a r s u i t o f e r e c t i v i t a t . a c i o n e o t h e r e s a t i v e l c h a f e a p h

o s . h a t e m u j m e l l n i q u e f i t n e s s o f v a t i f o l f o f m i n . h y . t o s e n s o r q u a e t e r i n n e s t i e f o p e r t i e s o f m a n i m p o r t a n t b i o m o j e c u j e s w a t e r e d p a r a c t a s a n d c . d a l p a r a s e i s i v i n i s i t s r e a t v e . s a t i l i t y i n b i o c h e m i c a l r e a c t i o n s i 2 i t e a t o l s i d a l l o t a i f e l i n e j e . i o . e s a t i v i t y e t h e r t w o u i d e n o p o i a n o n i s i t h i s w o u l d . f a s i c a - l y a r f e c t a l l r e a c t i o n s i h a t i n v o l v e y i c t e i o l d e s t o y r e s o l . a i . i n o x i g e l o r l i t o s e l i t h a t i s m o s t o f 2 3 w h a t a f e a c i d s a n a b a s e s ? i 3 i c c h 3 m e c o n i . s a t e a c i a c c h 3 n c o m l u s a t e i a s e # 3 n o e o d i c c o n t u s a t e a c i o .

H s . c h 2 - m e c o n s u s a t e h a s e + h y - c h 2 - m e c o n s u s a t e a t . a c i o h e m e - m e c o n s u s a t e h a s e # o r c h 2 - c o o h c o n s u r a t e a . i a # m e c h 2 - m e c o n s u r a t e . a s m e c h 2 - c o o h c o n s u s a t e a s h a m e c h 2 - c o o h c o n s u s a t e a c i a h a c a t r i b u t e n . a c o n s t a n t e s a b a t o t h e c o n c e n t r a t i o n o r t h e r o u l e s o r t h e a b e t i s = s o c i a t a d d i v i d e d b y t . e

c o . c e n t r a t i o n o . t h e u n d e r s s c i a t e a c i a f o r m e c h 2 - m e c o n s u r a t e o r a u t i v e o r a u t i v a t i v e a - s c r i p t i o n . f h o m l u c a c i a h a c i s s o c i a t e s t h a p r o f e n i o n c c y t h e r o p e r t o r a m o r e t u l e l i a h

a s . o f a p o r a r e s i o n n a . n o n p o r a r e s o n o c a l l e a t o u n o r a c i a o r y s t h a t c h e a d a t e o a u r e r .

cei luaitrav es ofka hno mānemost obinn Ati sjaqt it pleshty rā sie sšoxy ānā ān, tēcon cen tfaui Onof theaco xy forlbrt hēab nhrmanr mglorj nanc, easesl iśsoxygncan le lollfctau Sij n gthēōs efVēabre a thinslā, ficutiē-1, 2. IpfēqI hēm, glōbi btrēis ubu nātonfisi tj Onis22wā, hē tē alce hēlltūthē q dānsby thēc lai f rhes iCkē tēcei Im utātiōarf ecstūh bēch āi n sōtie fetlū spōlo, ysolls-Ofj bta sro mā tēta nēllm Oqōisānēā, hēreiatū Vbxyg tū afā hāticēuI Oxy ovyēl-Otē, āke nbyuere ā iCēil sēfūmēatēf hāhē, 154 B eca Usa tēpē opē ē i psicēle nēuēd ise āSe afēir onīC aqūā., ēllī cōsō hēcēlsw it hēfētalhār tē ro ducē atōhēi pōlē r cō tēf hāmpā i rēd oxyē., cēu vērys ysēm sō hēcry sta iū n ēror mhan gēabēca, sōxyēfēn tē fedlū r tē qvēr s hē p rāns rōf m n gē oxyē.,

MO, IO hān tōolY hē MO gōisānc hēpērsāns wēr, sō i Hōwōy ēExr āt-llēprō tēinsfō nēā i sē? I t bēkē r p O tē, nēvē, hēllm sōnā cāōp z i f vōntēdēd tōā i nēā i n thēst ructū rāi hē ēgr i t, rō f hēs u cē I I u I a r r g ā n i ē s, āp O t ē r nē l v ē j., ēll. Oūā tē tē tēf bēc āu scētis mōfē ēntū ēlth ē tīs sll ē. sū. hāsāvēr m lē tēso f tēnōuēlōs ēwīth is āē V. C. ēā āi t i ngōulī sāpō cēss wēr tēyānī s hū yōnī (sā itis i sē d tōf-āul, hēso, ubi i t yōrā p r o tē i nūltā i i t cōm ēsōtōr s., I u t i o nālic ān b cē ēntū f u sē ā t hēsā, fēllm s i o n d i p ōetōnā s w i t h t hē f a t ē r i n t hē sōlūtiōn w i c h i e q v ē s r e s w ā ē f a d i h b i e t h y d.,

at the p r o c e s s i n o f r a s i d c a d n s b e q n d i t i t a c u s e t e e n f o t ē i m b o p c u s ā n d i e v e . o n t h o r u s i e l t t e i n a n n o c i a t i o n i e n t a n i a r g n e m e n .

SM, kēsomep-ō-ēi nspōrts Oūllēth ānothērs āp, tōtēi nqā t hōpēnāshly pōi ārā mī nācā cōnt, ēsurrāc ē-i-ī tē mōrēso Iū b i ētānōr w i t h o f e n y ā f o p h o i c o l e , o n t e s u f f ā c e i h o l l e n i .

et, ēiv ēr ēusysnēp, o t i e n n e i v e i h e m o t o s e h i z z e n s i j n t i e h o m o s e n t ē e a t s o o x s t o s e a i t t i t u t i n h o k e n e i s a n ā , u c i e l c e n t r i , s u s t e s t h e p e n g t a n t a u s y o o o x g r a n a c o l i c e t h e p e l i e t .

iC, Cōn tāi p t h e m i t t c h o p d r i a l ā n o p e r o x i s o l l s ā l k i m i t o c h o l d , i a r ā v e o v e f i a , h i n s e d n e n ā h i o l c a r , c t e n s i c i o t e r t e c h i q l e s , s u t h g s u c h s e f a d e h e t t e r i f u s a t i o n w d u ā n v e t o b e u s e j c o , e r ā n a t e n t i y c o n s t e n t e s i t e s i n v n e a n c o o k i n e l l e a t a p e u . u ā u , m o r e t i a n n o u s n t o a e r ā n t t e l e p r o t e i n a r t o f t h e m e a p g , p h o n ā i s e l s e s t a v e b e e p l i n k e ā t o t h e i m m u n s y s t e m .

Sh, I i e v ē a t p a t t e r I i o n p r i t a n s t r a v e l i n d i y

MP, S Y s t e m b l u n d e O i y m p h o C Y e s a n d e v e n t y a l y ā r i v ē a t t e t e r l o u . t i s s l e w h e r e t h e y b e a n t o u r a n s o f t h e n o I m a l e f O n p t o t e i n t o t h a p r o f m a l o l e t o i w h i e p h e r t a r s .

FO, g s e p e i c p r e d s p o s i t i o n t l a q u e s c a r a t e , h ā t a ' O f e w ā l i m i t a u f t e a i s t a s e m e a i s e ā s e m i t b e s t ā r t e a v i s e . t i n ā p f i o n t a t a i f e a y ā s t e a n t e f e a c o l l f o m t i o l d . n p s c i u t e p r o t e i n e f r i c i e n c y a t i o s a r a n i t l a r y m ā s u r . m i n o f u e s s e n t i a l a h i n a c i a c o n t e l i t o f a s i v e n t y r e o p r o t ē i n i z e s s i a v e t h e n i g h t s t e r .

13. The g r a n n o a c i d s t h a n y s t r e c o n s u m e a i n t e l i e t b e ā u s e t h e b o p c o n n o t s y n t h e s i z e u e m i n s i f i c i e n t q u ā n t i t e s i n a R e d S o n s f o r t r e a n t i g e n e t j c h y m o d i f i e d f o s i n c l u e i .

C

UP, āfēpōtonāēqI āt p h i o y h o t h e n C l Δ r e x - I g o U p a n t - S i f C r a i n d f b o x y l g r o Δ u e d p i n t o l f d t o a r t f X y i a t e o n e d f t e a m i n o g r o u p s i p r i m a f i l Δ y e r - O t t i t e l a d h e o t h e i m

i n , g i o u r i g n i X t l r e o f u e p l o r o q u e ā n d e p r o t o n a t e d r o m s l e z i t e l r e f e r s t o t e r o m i h w i c h , O t h c a n o x l e r o l i s a r e a e p r o n a t e d a n a t o t i a n n o s r o u e s p i o t o n a t e a t p o s e z i t h e e r e f e . S t . t h e r o m i n l i c h o t h e f o X y i g f o u p , i f e a e p r o t i n a t e d a h a b t h a m i n o s r o u p s p r o t i n t e a t e t o g e l z a t t p h i t h e c h . r f e d , r o p s a r e t h e n o m e r i t a l n s t o n ā i n p l a n a t h e p r o t o n a t e a g l a n a i .

O g r o u p n a r s i n j i e s i v i n g z e f o n e t a r g e i

sat p h a r i t h h e c a r i o x i a t e r o u p o n t h e c t e r m i n a l l e u c i n o s i v i n s a t e c r a f

Δ T e - a r e d g r o u p Δ e e s a e s p h a Δ t e a d t i o n o f

Δ T e - a r e d g r o u p Δ e e s a e s p h a Δ t e a d t i o n o f

Δ T e - a r e d g r o u p Δ e e s a e s p h a Δ t e a d t i o n o f

Δ T e - a r e d g r o u p Δ e e s a e s p h a Δ t e a d t i o n o f

Δ T e - a r e d g r o u p Δ e e s a e s p h a Δ t e a d t i o n o f

UP, āfēpōtonāēqI āt p h i o y h o t h e n C l Δ r e x - I g o U p a n t - S i f C r a i n d f b o x y l g r o Δ u e d p i n t o l f d t o a r t f X y i a t e o n e d f t e a m i n o g r o u p s i p r i m a f i l Δ y e r - O t t i t e l a d h e o t h e i m

i n , g i o u r i g n i X t l r e o f u e p l o r o q u e ā n d e p r o t o n a t e d r o m s l e z i t e l r e f e r s t o t e r o m i h w i c h , O t h c a n o x l e r o l i s a r e a e p r o n a t e d a n a t o t i a n n o s r o u e s p i o t o n a t e a t p o s e z i t h e e r e f e . S t . t h e r o m i n l i c h o t h e f o X y i g f o u p , i f e a e p r o t i n a t e d a h a b t h a m i n o s r o u p s p r o t i n t e a t e t o g e l z a t t p h i t h e c h . r f e d , r o p s a r e t h e n o m e r i t a l n s t o n ā i n p l a n a t h e p r o t o n a t e a g l a n a i .

O g r o u p n a r s i n j i e s i v i n g z e f o n e t a r g e i

sat p h a r i t h h e c a r i o x i a t e r o u p o n t h e c t e r m i n a l l e u c i n o s i v i n s a t e c r a f

Δ T e - a r e d g r o u p Δ e e s a e s p h a Δ t e a d t i o n o f

Δ T e - a r e d g r o u p Δ e e s a e s p h a Δ t e a d t i o n o f

Δ T e - a r e d g r o u p Δ e e s a e s p h a Δ t e a d t i o n o f

Δ T e - a r e d g r o u p Δ e e s a e s p h a Δ t e a d t i o n o f

Δ T e - a r e d g r o u p Δ e e s a e s p h a Δ t e a d t i o n o f

Δ T e - a r e d g r o u p Δ e e s a e s p h a Δ t e a d t i o n o f

Δ T e - a r e d g r o u p Δ e e s a e s p h a Δ t e a d t i o n o f

Δ T e - a r e d g r o u p Δ e e s a e s p h a Δ t e a d t i o n o f

Δ T e - a r e d g r o u p Δ e e s a e s p h a Δ t e a d t i o n o f

thēbno m alprionp, Ot ēi n i n c o l l s , t i e a i s e a s e c a l l s e d b y p r i o n s i c a i l e a n v i p e s o n s i f o p h e c e p h i o r a t e y p l o r e c o m m n l y m i d c o w d i s e a s i n s h e p t h e j i s e a s i s c h i e a s . f a p e r i n h i d i s i t i s c a l l e d a k o b , c f e u z f e i d a i s d e e s o i h e n o r m a u f o r m o f t e r e p r i o n p f o t e i m a s a h i s e a m e l i x c o t e n t c o l p a r t i d .

h e t h e a n d m a i o r e h a s a n i n c r e a s e a s e a 4 0 7 w a t r i s a b o u d t e m a r s t u c a r e c o r o .

e n s o j i n h o c o n t a j t h e r e g r o u p i n u a r e d y e n p i k i n s s e c o n d a r y s t i u t y t e i s p r i m a f i n y m e l i x i j i e r e r e s e s i s a t e t r a m .

n w a j e m y o l o b i n i s h o t o t e r o x i g e n j i n i s t o e m .

g l i n i s c o o p e r a t i v e h i m n c o o p e r a t i v e t o m y o g l o b i n a l t i e c u c i a , r e s i d u e s a r t h i s t a i l s i n s o m p r o t e j i s t a d i h s h e s t e r e v o f o r g a n i z a t i o n i s t e r i . f y h a n o s o b i n i s i s q u a t e f a i y 4 3 t h e r l i c t .

n o , h e n o g l o b i n i s o x y e n t i a n s e o r t j i s s i s d i d l y i n i n s c u l y t r e f i e c t s t e r e a c t i n a t e r i n b i n d a s i n t o o x y e n t e o m p a r t j v e r y h i s e r e s s u r e s a n i t i e a s e o x y g e n a t y o w e r e s s u r e s i t e r .

n c i o n o p i o s t o b i n i s o b s e r v e s t o r a e a s a f e s u l t i t i s e a s i n s a t u l a e a w i t h o y s e l a t o . p r e s s u r e s a s s h o w n i t s h y e n o i c b i n d i

c u , v e l 4 a i t t e r p e s e n c e o f f i d a l a c o l b o t h o r m a c h i n i n a t o h e m o s i o n i n t , e o x y e n j i n a i n s c a p a t e y o n e n d g l o b i n a e t r e a s e i s i n t h e a s e n c e o f z o n e s p h o s i .

e a t e r e b i n i n s a f o x y e n b y h e m p s o l j i n f e s e l l e s t r a t o m y o g l o b i n c l a r a t e r i z e a b y p l a c o r e . o p e r a t i v i t y i 2 3 n p h o s p h o s i n t e r a t e b i n a s a t e n t e n t e r o f t h e r e l l o s i o b i n i o r e c y l e i n c r e a s e s c o o p e r a .

i v , t y s t a p i i z e s t h e o x y o n f i l t a t i o n p h e m o g i o i n a n i m o u n t e s t e b i n a i n s o t x y e n o u a t i t e d n e i s i y h e r e a s e d i n t h e c a r i n a r i s 4 0 f e t a l h e m o g l o b i n .

i n , s o x y e n m o r e s t r o n g l y b i n d s t h e m o g i o b i n s e e i s u r e a 2 3 4 i n s t i d i n g l a j i n a c h a i n s p e r i c e b .

a s , f i n i n e a s t e o x y e n a t e d h e m o s i o b i n i s a p l a c e r a c i a c n a s h i s p r o p a r t a p p o x y e n a .

t h e m o s i o n i n m o t e r i f o f d o x y e n e a d e a t h e m o g l o b i n i n i n d s m o r e s t r o n g l y o b i n a d e , o x y e n a t e a h e m o s i o n i n t e b i n a n g o m e c a n a f c o b . o n e m o s i o b i n f a v o r s t h e c a n a t e r m i n a t e r m i n a t e r u c l i e t o t h e a o x y e n a t e a t o m o r h e m o s i o b i n i n t h e p r i m a

y f , w i t h o u t r i e n a s t e a s o n i s s a r t e r s a i o f t h e a f i n i t i o n o f t h e w h i c h s p e c h o s h t a l l f t h . t e l e s t o r s i n a n g o n y a r o g e n o n s y n e n o s o b i n i n f e t e r e p h i m a r y a .

t o , i n t e r o f e r e c t o t h e p h a n g e s t o u , d b e t h e o p o s i t e o f t h e a t e m a n y o b s e r v a t i o n e s , e s o n s o f h e m o g l o b i n t o c h a l e s i n p h i s t i c e n t r a p o i n t w i t h t h e p h i c r e a s e s t h e h i a r o n i c o l c o n c e n t r a t i o n .

e c e a s e s a n d i c e v e r g a s o t h e c h a n g e o f a n i s t a i n e t o a s e r i n e .

t i e h a i n e m o y s i o s i n t i v e l y c h a r a c t e r i s t i c a .

h a c o u n d a v e i t e r a c t e d w i t h t h e t h e r e a l t e w e i s a j t h i c a s t o n e a n d s h a n a , n s i s e a s i e r t h a n i s i n a n s i p e r s o n s w i t h s i c k

ar ysruurē of f o t e j n s ? 15 1 share) .
 O L , b j i t y a n d t e f e o b i o n o g i c a n u i c t j O k S t a t i c , s i t u l t i r a
 v f u s . Y n a m i c , c a l a i y t i o i t r i e a n f i b o f t h e a n i d e p a n s -
 s t h e y r o t g e a t o l l t u e a l e i f b o n T h e a l l e s e s a r e b o t h a e r i n e
 c i a s e r j o w . n t h e t o p I g n e s t o u I a s e o v e f i a p p i f f u c h t a u t h e l
 a n t o n y . S i o u r o r j n e c o m t a t S u e n n h y t h e o t r e p l 1 7 h o u i g i -
 a l o m m o n O n r e r e t i t i v e i r r e s u l a n t y f o u p a i p a n t i l a r , a l l e l S h e
 e t s I A m i s a u i g n m e n t o c c u . S b e t t e f s t r a n a s o f t h e ■ S t e e t a
 S a n S h e s i a e t o t o p p o u W a r a n s A k e v e r .
 e t . n i S a f e ² , o n o r a b I y r e p t i a e w h e t e .
 i f . C t i o n C h a n e S b a r o u t i s o j i t e r e a r e u o . i l d - t h o s e t i a
 t o l l a , n p r o i j n a n a u s e t e t h a t o n o t i s t e f i s u r e a o f o r t x a
 n p e s h 9 T h e a l t h i s i s t e f u l l y e x t e n d a e s a n a t s i n d r o g .

n h l d s a f e p a n a i e i t o u e p r o t e i n f i b e n t h e I P r e a t e d h e l
 t s . n j t u r e i a i m o s t u n y e x t e n d a n a n i t s h y a r O l e n o n a s i f e
 p e r t h a i t a r t .
 T h e p r o t e i n a i e r i 2 o i n t a a l l i e t h e j a n u i t y e h e n p a n e b t i e t -
 e k a e y t h e n a 2 i t h e g e o n e t r y o r t e p o u n e f e s i d e i s s u t h a t a i t d .
 e s . o t f i t j n o t h e a n e i x b u i t a e s f i t e x a c t I y r o j a r e v e r s e u f
 n s e e f a g h e 2 2 i g , Y c i f i s t e t e o n y r e i a u s m a i e n o u g .
 t o . i t a t e r u i a i p o i n t s i n t h e c . I n a e n t i p i e h e l i y 2 3 i t h e p a n i
 c i p a i c o m m o n e n o r w o i t i s t e p r o t e i n e r a t i n y w i c h y s i c i a . S i c e
 x a m p l e o f ■ h e i c a i s t r u c t u r e t h e p r i m a r i C o m p o n e n t o f S i n e
 i s t h e p r o t e i n f i b r o j o i n t h i s i s a c i a s s i e f x a m p l e o r t h e e i s t r u c
 t u r e t h e s t a t e .
 t j . s o m e w h a t e r a n o v e r s i m i l i t a c a l l o b b e i t i s p u n a m e n t a l y -
 a n d i 2 d w o t w i c h c o f s i s t s a l r e i y o t h e r o t e i n k e f a u t h
 s h r i n k s b e c a u s e o f a t s n e y C a l l o f f o .

12. i O n i c o n s t i t u t e h a n t e n S t r i n k s i c o n s i s t s i a n e l x o f -
 f e r s . t e i n f i b o i n t h i c h i s t h e f l u i d y e h e d e s h e e t c o n r o f f a t i o
 n y w i t h a . I e s t e n d e n c y t o s t e t h o i s h i n k a t w h a t a n e s
 a y b o u t h e r e t h o d y n a m i c o f p r o t e i n ²⁵ (i) b a c k r o p e t m o p a s .

i n . O i v i n g h e c a .
 d N , g o u p s o f t e l e t j e c h a i n i c 2 s i a e o d a i n h m o l d s i n y o l i n s a
 n y p o s s i b i e h y a t e s e . b o n d d o n o r o p a c e p t o r s o n t h e s i a t e l a
 i n s : > h y a r p h o b i c a r t e f a c t i o n s i n v o i v a n s e t e n o n p o i a
 p r o u s o n t h e p r o t e i n s . d e i t t r o s t a t j C i n e r a c t j o n s i m o u
 v i n e a l y c h a r a c t e r . o l l e s o n t h e p r o t e i n : < s > n e t a u l i s a p i o n i
 n o i v i l g c o o r a i n a i o n s o n i s b e t e e n s i a e c h a i n s a l l a l l e t a q u o n 2

13. a s i l i z a t i o n o r a p c o n t r o l l a t i o n d e p e n s o n l o Y i n s o f e " I g s i e
 s a r a e s o f t h e r i n t h e i n e r a c t i o n i n v i v e a n i t s m a i j u l l e d i e
 i n t e r a c t i o n s , t h e l a n C o m .

o n . n t o f e n e r g i c h a n g e s i s e n t i a l p i c o w i t h r e l a t i v e l y s m a l l e n t
 r o p y - h a n s e s i n t h e a s e o f m o r e . U l e s s i a r e q u a s i r o t e j n s t e n d i n g e
 r o f p o s s i b i e c o n t r o l l a t i o n p h a s e s t h e e n t r o p y (p a r t o f n e r s y d a -
 s e s m u c h a n o t e i n p . o r d i n u m i s i s p a t i c u a l l y r e l e a n e c l o s e o f y
 d o n o b a c i n t e r a c t i o n . n 4 o s p a t i s t e t e r t a d y s t u t e p r e s e n t o t
 e i n s ? 7 s e e h a r e a 2 r o r a n y d i l l g t h o t h a t a i s h a t e 
 t h e r e i x s t e a n a r i s t . C a r e o i s e f i s u r e a 1 a f o r a n y a r o s e
 n o l i a t a t i s p a r t o f . t a a r y s t i c t u r e s i a e l p i n y a r o e n e r o n

i n s l 2 8 S e f i g u r e a 1 3 r o t e c t . s l a t i c i n e r a C t i o n s u . h
 a s i g h t e s e t h b e t w e e n . h e s i d e h a i n s o f i y s i n p a n d a s t a r t a t e
 2 9 s e e e i g u r e a n d r o r d p x a m p l e o f a a j S u i p a e s o n d j 3 0 s e e
 F i s u r e h i 3 f o a e .

20. I e o n y d r o p h o b i c b o n d s i 3 1 a c h a p e r o n e i s a p r o t e i n t h a t a i d s
 a l o u e r p r o t e i n i n r o , a n e o r r e c t i v a n a k e p s i t e f o n a s s o c i a t i o n e
 h . t h e p r o t e i n s i e f o r e i t h a s t a c l e a i t s i a n b i t t l r e f o r m
 a i o n n i a l p r o t e i n .

f e . s t o t e p o s i t i o n o f i n u s a l e t c o v a l e n t o n a i n s E x a m p l e s i
 n e j u a e a n n i s o m e r a n a o p t i c a j i s o m e b l o c k o f o n a t i o n r e f e r s t o
 t h e r o s i t . O i n g l o s s i o u s i n s i l l e a l l e t o r o t i t i o n a t o h a s i l e .
 e b o n d s . A n e x a m p l e i s t h e d i f f e r e n c e b e t w e e n t h e c h i s t e a n d s a d s
 s e f e a o n e . I n a t o n s o r e t r a n s a 3 1 F i v e p o s s i b i e f e a t u r e s i j t p
 o s s i b l e p b .

e i . C o n i s u r a t i o n s a n d c o n f o r m a t i o n s (1) .  h o u s a n y o n e o f t h a t i n o a c t i s i s p o s s i b i e a t e a c h p o s i t i o n
 o n t y o r e i s i s e d a s d i c t a t e a n t .

e g . n e t i a t c o d e s r o t h a t p r o t e i n (2 > i t h e r a n b o a r l l a m i n o r c i
 d e o u a n e u s e a t e a q u o s i t i o n e x c e p t o r a l y e i n e) , b u t o l l y
 l a a n y n o a c i a r .

U S . a l e s t h e r e t a d e s t o p i s p a p p r s , t h a o n l y a n d a n a f r a n s e p e n t
 S t r e o s s e r v e d e n e r g y s t o r . i s m o r e s t a b i l e a n d s t h e c o n s e r v a t i v e f o u
 n i a p r o t e i n s (a r t h e n s e s a l a c a n t e n o r e i t a l i y a k e n a n n a n e .
 I O . t o t o o i n b u t s o l l e a n l e s a r e n o t p o s s i b l e b e c a u s e o f s t e r i c h
 i n a r a n c e s . N e j e s t i a q u e s t i o n a l i y a n o w e , m a n o t h a v e s t
 a n i z i n g t h e q u a n t i t y o f s u c h s t a b o s a n t h e n e i x c o r t h .

P r . i n a s t r u c t u r e t e r m i n e s a n o f i m u n t e r t a l l s e l l e t e r , a c c o r d
 i n s t o t h e i s c o l l a t a i r o t h e e l e n e t j c o . e l i 3 4 t e c h n i c a l i y o l o
 r a g e n a s t h a t e f a r y s t r u c t u r e l a s e i t h a s m o j u l i p i e r o y y ,
 e p t a d e c i a i t s . h o w e r e n i l l s t a i s C u s s i o n s o r y u l t e r m a l y s t r
 u c t u r e i n o u e s t h u n e s o f e i o p u d a r p r o t e i n s o f f i b r o u s o n e s i i t e c
 O l i a s e l) m a n y s u e l i t i s t s c o s i a e r t h e c o l l a s t i t r a p e r e i x t h y
 a r t x a l t e o r t h . C o l l a p y s t r u c t u r e h o s e 3 4 s i i l h a o e c y i s u i a t e r o
 y a c r y a n n a s e j e l e c t . o p h o f e s i s i w i t h i s I P A G e t h e h a s e
 a n a n o e d i f f e r e n c e s o f p r o t e i n s t r e i i n i n t e d s o . t a t h e o n
 s t a r a l l e g e t e r m i n i n s t e l l i s a t i o n s t h e s i z e o f t h e p r o t e i n s i s
 D s b i n s t o u e p r o , e i n i n a c o l l a n t i a t i o o f p l a s s i s t e r . t a m o m e n t o
 a n l e o u s t h e r o t e j n w i t h e s q u a n t i t y e s a n a p l u s i a n o d a n o
 n e o j i s h a r e . T h i s c h a r a c t e r i s t i c a f e e l i m i t a t e j 3 6 i n a r o u a
 c r y s t a l l e s e i n s e a r o f e t r i l f a t i o n c h i o m a t h e r a p h) t h e l a r g e r p .

O t . i n s c a n t l a v e i a r o u n t h e l a s i t h e r e s . i d v i n s a s h o r t e f p a r t n e o
 t f a v e a n e l u t i n g a s t e r t i t h e e t t r o p h o r t S i s t h e p
 f o t e i l s a t e f o r e a t , s o m e u s h e n a t e r s o t t e y t r a v e l i n o f e s i o
 w i y a t h e v l r t i s s e f i c a u s e t h e r e i s n o o f f i c t i o n l 3 7 T h e m i s
 3 7) o d a u t h j s i a n o . e d e t e r m i n e t e r . i m a f y s t r u c t u r e o r a p
 p o t e i n 2 3 s i n e t h e n g l e s . a d a t i l l W i l l i a m s i w e i d e n t i t o r t e n e
 t e r n i n y s i n a t s e f i r s t c y c l e s o a n n a s e p a r a t e x p e r i m e n t s o -
 n e c e s s . i y i 3 9 i t m i s t t e l l y o u i a t h o f o t e i l w a s p u t e o r i f t h
 e f e w e r e s u n d e r t s .

4 . I t h e a m o u n t o f e a r a n d a s e n t i m u s t e x a c t i m a t e r e a m o u n t o f e
 t e l m a n i t h . f i r s t r e a c t i o n i n t h e r e i s t o o i t e r e i n a n e s e n e

a t a j r e a d y n a m i c t h e a n t e . C o n t r o l l a t i o n o f P s o i b i t e r o t e i n e f f
 e i e n c y a t i o n i s h a n t i e f a y m e a s u r e m t h t i f t i e s s .

n t . a i a m i n o a c i a c o n t e l t o r a s i v e n t y p e o f p r o t e i n l 2 i e s s . a v
 e t e r n i g h e s t e r l e s i t h e a m i b o a c i a s t h a t i l l s t h e o n s u l e a i n t h
 e i t . b e c a u s e t h e o r a c a n b o i s i t l e s i z e e n i n s u f f i c i e n t e u a n t
 i t a e s l a r . a s o n s r o r c r e a t i n g t h e t a c a u p l o d i f i e a f o o a s i -
 c u l e i n c l e s i l l e s t h e i f r o t e i n c o l l . n e j l e r e s i n g t e i f s -
 e r n i t h i n d e a s i n a t e i f i s i a l r e t e i n s e c t s o m e t e r r e s i s t a -
 d e c f e a s i n g t h e t e f a t o r e s t i c i a d s t e r . w t h e l 4 . a w h a t s h e s .